GenCore version 5.1.5 Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - nucleic search, using frame_plus_p2n model

May 31, 2003, 18:52:35; Search time 1352 Seconds (without alignments) 2455.677 Million cell updates/sec Run on:

1 MAHAGRIGYDNREIVMKYIH........HTWIQDNGGWYGASGDVSLG 205 US-09-375-514-22 1104 Title: Perfect score: Sequence:

0.5 7.0 7.0 Xgapop 10.0 , Xgapext Ygapop 10.0 , Ygapext Fgapop 6.0 , Fgapext Delop 6.0 , Delext BLOSUM62 Scoring table:

16154066 seqs, 8097743376 residues Searched:

60362 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 10 Maximum DB seq length: 40

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 500 summaries

Database :

em_gss_mus:* em_gss_other:* em_gss_hum:* em_gss_inv:* em_gss_mam:* em_gss_vrt:* em_gss_fun:* em_estba:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

t the total score distribution.	SUMMARIES	Descript	999756 S893e04.	24 AISIIU24 TFU3AU3. 27 AIF98327 +*63<12	180 AI202980 qr32e08.	006 AI628006 £y21d06.	10 AI422710 tf29d12.	1450 BMOLO450 6036.	MODEL 1972 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	700 Al039224 Homo se	AZ782295 2M002	1288 BE/29288 BUIDS. 32 ATG10530 +20540	A1131930 uc33q07	1681 AZ832681 ZMOIII	744 BE395/44 BULST	1439 BG709439 602673	1387 AZ780387 2M0017	AZ784258 ZM0026	135 415291(1529) 11 AI362411 av92e03	02 AI612902 ty33d0	AI761879 wg68c11	08 ALULS468 OVS46LL 61 ATROU161 +r235DR	483 AZ982483 2M026	272 AW250272 282118	54 AA999756 OS93e04	81 AI677781 WC80C06	555 BL/62555 60304E	158 BI833158 603090	514 BI833514 603088	654 BI829654 603079	97 AI589397 tr61h11	979 888 AZ477888 1M0297	266 BH792266 SALK_0	oi AA98/401 og8/g06 25 AI357425 gu01c09	358 BI829358 603079	75 AA869375 vg07c10	02 AL303182 LQ30N1U 178 A7373178 1M0133	536 BE867536 601443	39 AI024239 ov71g02	470 AZ791470 2M0041	AU3P AL478544 T. bru 65	03 A12/3863 QUUSALU 14 A1287914 QU13CO6	57 AI364457 qw38e11	03 AA625403 af69a08.r	481 BI765481 603050	539 AZ313539 IMOU30KU 894 BG973894 60284352	4 T17564 mps v232	48 AA034448 ZK23a08	54 AA8/8864 OI849U5	41282/21 GT65CIU
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## os93e04.sl NCI_CGAP_GC3 Homo sapiens cDNA clone IMAGE:1612926 3'similar to TR:062084 Q62084 PHOSPHOLIPASE C NEIGHBORING;, mRNA sequence. AA999756

LOCUS DEFINITION AA999756/c

RESULT 1

EST 05-JUN-1998

linear

mRNA

31 bp

AA999756.1 GI:3190311 Homo sapiens numan. SOURCE ORGANISM ACCESSION VERSION KEYWORDS

REFERENCE AUTHORS TITLE

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 31)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),

JOURNAL

Tumor Gene Index Unpublished (1997)

Contact: Robert Strausberg, Ph.D.
Maral: cgapbs -femil..nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael
Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: ww-bio.llnl.gov/bbrp/image/image.html

Trace considered overall poor quality Seq primer: -40ml3 fwd. ET from Amersham High quality sequence stop: 1. Location/Qualifiers

source FEATURES

/ulliance | Interpretation | Interpretat /organism="Homo sapiens" /db_xref="taxon:9606" 16 g constructed by

BASE COUNT

Conservative: Mismatches: Indels: Length: Matches: 1.09e+0544.00 88.89% 77.78% 3.99% Similarity: Percent Similarity: Alignment Scores: Query Match: Pred. No.: Best Local

US-09-375-514-22 (1-205) x AA999756 (1-31)

AI698327 37 bp mRNA linear EST 16-DEC-1999 tx63c12.x1 NCI_CGAP_Ut1 Homo sapiens cDNA clone IMAGE:2274262 3' similar to TR:Q41120 Q41120 HYDROXYPROLINE-RICH GLYCOPROTEIN ;,

AI698327.1 GI:4986227

Homo sapiens

ORGANISM

AUTHORS TITLE

REFERENCE

human.

mRNA sequence. AI698327

ACCESSION

KEYWORDS

SOURCE

VERSION

AI698327/c DEFINITION

LOCUS

RESULT 3

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24000

Conservative: Mismatches:

42.00 77.78% 77.78% 3.80%

Best Local Similarity: Percent Similarity:

Query Match:

Alignment Scores:

Pred. No.:

Score:

Indels:

JS-09-375-514-22 (1-205) x AIB11024 (1-34) 39 ProProGlyAlaAlaProAlaProGly 47 CCCCCGGAAAAGCCCCCCCCCGGGA 30

Length: Matches:

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 37) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP),

39 ProProGlyAlaAlaProAlaProGly 47 30 CCGCCCGGACCCTCTCCTGCGCCCGGG 4 ŏ

LOCUS DEFINITION RESULT 2 AI811024

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Email: Crapbs remain.in. gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-GGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/db_xref="taxon:9606"
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High quality sequence stop: 1.
Location/Qualifiers
                                                       GI:5397590
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Unpublished (1997)
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AI811024
AI811024.1
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ORIGIN
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TITLE
ACCESSION
                                                                                                        KEYWORDS
SOURCE
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JOURNAL COMMENT

source

FEATURES

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40 bp mRNA linear EST 07-MAR-2000 ty21406.x1 NCI_CGAP_Ut3 Homo sapiens cDNA clone IMAGE:2279723 3' similar to TR:043687 043687 EXTENSIN-LIKE PROTEIN; contains element MSR1 repetitive element; mRNA sequence.
A1628006
A1658006.1 GI:4664806
EST.
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Email: cgapbs-rémail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.

Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: Life Technologies, Inc.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/lmage.html
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                                                                                                                                                                                                                                                                                                             /tissue_type="pooled_germ cell tumors" /lab_host="DH10B"
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Insert Length: 1838 Std Error: 0.00
                                                                                                       Trace considered overall poor quality
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Matches:
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/clone="IMAGE:1942598"
/clone_lib="NCI_CGAP_GC6"
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                                                                                                                                                                                                                          /organism="Homo sapiens"
                                                                                                                          Seq primer: -400P from Gibco
High quality sequence stop: 1.
Location/Qualifiers
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Seq primer: -400P from Gibco
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Unpublished (1997)
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KEYWORDS
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                               Unpublishe Index (1997)
Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
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94 bp mRNA linear EST 14-OCT-1998
9732608.xl NCI_CGAP_GC6 Homo sapiens cDNA clone IMAGE:1942598 3'
similar to TR:Q08805 Q08805 SALIVARY PROLINE-RICH PROTEIN L
CONTAINS element MSR1 repetitive element; " mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone_lib="NCI_CGAP_Utl"
/clone_lib="NCI_CGAP_Utl"
/clone_lib="NCI_CGAP_Utl"
/clone_lib="NCI_CGAP_Utl"
adenocarcinoma, 7 pooled tumors"
/lab_host="DHIOB"
/note="Organ: uterus; Vector: pCMV-SPORT6; Site_l: SalI;
Site_l: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.75 kb. Life Technologies catalog #:
11538-014"
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DNA Sequencing by: Washington University Genome Sequencing Center
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Sukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 34)
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Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism-"Homo sapiens"
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AI202980.1 GI:3755586
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Unpublished (1997)
                   Tumor Gene Index
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Query Match:
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Pred. No.:

Score:

BASE COUNT ORIGIN

AI202980 OCUS

ð g SEFINITION

ACCESSION VERSION KEYWORDS SOURCE ORGANISM

REFERENCE AUTHORS JOURNAL

TITLE

COMMENT

31 0 0 0

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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1445310"
/clone="IMAGE:1445310"
/clone_lib="NIH MGC.41"
/tissue_type="amelanotic melanoma, cell line"
/tissue_type="amelanotic melanoma, cell line"
/tissue_type="amelanotic porpy; site_l: xhoI; site_2:
/lab.host="Dryn: skin; Vector: pOrpy; site_l: xhoI; site_2:
EcoRI; cDNA made by oligo-dT priming. Directionally cloned
into EcoRI/XhoI sites using the following 5' adaptor:
GGCAGGG(G). Library constructed by Ling Hong in the
laboratory of Gerald M. Rubin (University of California,
Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and
Superscript II RT (Life Technologies). Note: this is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BM010450 36 bp mRNA linear EST 30-OCT-2001 603631395F1 NIH_MGC_41 Homo sapiens cDNA clone IMAGE:5445310 5',
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mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 36)
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CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1925 row: m column: 23
High quality sequence stop: 36.
Location/Qualifiers
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
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Mismatches:
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BM010450.1 GI:16524804
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Best Local Similarity:
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31 bp mRNA linear EST 30-MAR-1999 tf29d12.xl NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2097623 3' similar to SW:CAl3_BOVIN P04258 COLLAGEN ALPHA 1(III) CHAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bonaldo, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                      /note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sal1; Site_2: Not1; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.45 kb. Life Technologies catalog #: 11541-018"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: David N. Louis, M.D., Myrna R. Rosenfeld M.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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NCI/NINDS-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NCI/NINDS-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NAI/ONAL Cancer Institute / National Institute of Neurological Disorders and Stroke, Brain Tumor Genome Anatomy Project (CGAP/BTGAP), Tumor Gene Index
Unpublished (1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima
                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2279723"
/clone_lib="NCI_CGAP_Ut3"
/tissue_type="poorly-differentiated endometrial
adenocarcinoma, 2 pooled tumors"
/lab_host="DHIOB"
                                                                                                                                                                                                                                                                                                                                                                                                                 0 0 0 0 0 0 0
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/db_xref="taxon:9606"
/clone="IMAGE:2097623"
/clone_lib="NCI_CGAP_Brn23"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-375-514-22 (1-205) x AI628006 (1-40)
High quality sequence stop: 1
                                         Location/Qualifiers
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A1422710.1 GI:4268641
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61.54%
53.85%
3.71%
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Best Local Similarity:
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AI422710/c
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                  EST 16-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: Sal1; Site_2: Not1; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.48 kb. Life Technologies catalog #: 1142.016"
                                                                             AI682252 37 bp mRNA linear EST 16-DEC-1990 tx02e12.xl NCI_CGAP_Ut4 Homo sapiens cDNA clone IMAGE:2268046 37 similar to TR:Q04118 Q04118 SALIVARY PROLINE-RICH GLYCOPROTEIN G1 PRECURSOR.; contains TAR1.b2 MSR1 repetitive element; mRNA
                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. I Chases 1 to 37. CASA http://www.ncbi.nlm.nih.gov/ncicgap. NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /tissue_type="serous papillary carcinoma, high grade, 2 pooled tumors" /lab_host="DH108"
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Matches:
36 ATGGGGGCGTCCCTCAGGGCCTGGCTCCTCCTCC 1
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2268046"
/clone_lib="NCI_CGAP_Ut4"
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Seq primer: -400P from Glbco
High quality sequence stop: 1.
Location/Qualifiers
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40.00
70.008
70.008
3.628
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Unpublished (1997)
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VERSION
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AI683252
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/db_xref="taxon:9666"
/db_xref="Taxon:9666"
/clone_lib="INHIE_MGC_41"
/clone_lib="NHIE_MGC_41"
/tissue_type="amelanotic melanoma, cell line"
/tissue_type="amelanotic melanoma, cell line"
/tab_host="bullulu (phage-resistant)"
/note="Organ: skin; Vector: pOTB7; Site_l: XhoI; Site_2:
ECORI; cDNA made by oligo-dr priming. Directionally cloned
into EcoRI/AnoI sites using the following 5' adaptor:
GGCACGAG(G). Library constructed by Ling Hong in the
laboratory of Gerald M. Rubin (University of California,
Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and
Superscript II RT (Life Technologies). Note: this is a
NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                            BM010792 36-0CT-2001
603629306F1 NIH_MGC_41 Homo sapiens CDNA clone IMAGE:5434872 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Tissue Procurement: DCTD/DTP
CDNA Library Preparation: Ling Hong/Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LINL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/Linl at:
http://image.llnl.gov
Plate: LLCM1912 row: k column: 01
High quality sequence stop: 36.
Location/Qualifiers
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Marazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 36)
MTH-MGC http://mgc.ncl.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
                                               22 7 36
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Matches:
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BM010792.1 GI:16525146
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                                                  Pred. No.:
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Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
     Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA sequence.
BE729288
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                                                                                                                        Homo sapiens (human)
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/clone="DKFZp(null)E191"
/clone_lib="(null) (synonym: hmcfl). Vector pSportl; host DH10B; sites Not1 + SalI"
/dev_stage="ault"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AZ782295
2M0022019F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0022019 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Toron (1) (Dases 1 to 27)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,

Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                            Homo sapiens mRNA; EST DKFZp(null)E191_r1 (from clone DKFZp(null)E191)
                                                                                                                                                                                                                                                                                                                                                    This close is available at the RZPD in Berlin
Please contact the RZPD: Ressourcenzentrum, Heubnerweg 6, 14059
Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de
                                                                                                                                                                                                                          Poustka A., Klein M., Mewes W., Gassenhuber J., Wiemann S.;
                                                                                                                                                                                                                                                     Submitted (12-MAR-1999) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                    Clone from S. Wiemann, sequenced by DKFZ within the cDNA sequencing consortium of the German Genome Project No s1 sequence available
                                                                                                                                                                                                                                                                        MIPS, Am Klopferspitz 18a D-82152 Martinsried, GERMANY
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Mismatches:
             12-WAR-1999 (Rel. 59, Created)
12-WAR-1999 (Rel. 59, Last updated, Version 1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6 A; 16 C; 9 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="breast cancer"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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                                                                                             EST; expressed sequence tag.
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40.00
88.89%
66.67%
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Unpublished (2000)
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gal electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil47321141gblAR129072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically competent E. coli XLLO-Gold (Stratagene) cells and selected for ampicillin resistance."
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 34)
                                                                                                                                                                                                                                                                                                                                                                                                                                                              /lab_host="E. Coli strain XLIO-Gold, Tl-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NIH WGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                             /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0022019"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
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Mismatches:
Indels:
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                                         0.00
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Matches:
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0022 row: 0 column: 19
Seg primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                        /organism="Mus musculus"
                                                                                                                                                                 High quality sequence stop: 27. Location/Qualifiers
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39.00
87.50%
75.00%
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us-09-375-514-22.rst

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Mus musculus
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similar to TR:056971 056971 ORF FOR OYERLAPPING PROTEIN: ;contains
MER22.b2 MSRI repetitive element;, mRNA sequence.
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CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP Clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                 /tissue_type="melanotic melanoma"
/tissue_type="melanotic melanoma"
/nab_bost="bull0B (phage-resistant)"
/note="organ: skin; vector: poTb]; Site_1: XhoI; Site_2:
ECORI; cDNA made by oligo-dT priming. Directionally
cloned into ECORIXAhoI sites using the following 5'
adaptor: GGCAGGG(G). Size-selected >500bp for average
insert size 1.8kb. Library constructed by Ling Hong in
the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
s 5 c 15 g 6 t
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 31)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
Bmail: cgapbs-remail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
           Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Tissue Procurement: ArCC/DorD/DrP
CDNA Library Preparation: Ling Hong/Rubin Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: image.llnl.gov
Plate: LLCM506 row: j column: 14.
Location/Qualiflers
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Mismatches:
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Matches:
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/organism-"Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:8831277"
/clone_lib="NIH_MGC_20"
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Unpublished (1997)
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AI912532/c
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FERSION
KEYWORDS
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AUTHORS
JOURNAL
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Eukaryotta; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 34)
Marrah, M. Hillier, L., Allaler, M., Bowles, M., Dietrich, N., Dubuque, T., Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptoe, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylle, T., Lennon, G., Soares, B., Wilson, R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EST 14-SEP-1998
                                                                                                                                                                                                                                                                                                           /note_"Organ: uterus; Vector: pCMV-SPORT6; Stte_1: SalI;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.85 kb. Life Technologies catalog #:
11539-2012" 0 g 3 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        uc33g07.rl Soares_mammary_gland_NbAMG Mus musculus cDNA clone
IMAGE:1399836 5' similar to TR:035054 035054 CPE-RECEPTOR. ;, mRNA
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                                                                                                                                                              /db_xref="taxon:9606"
/clone="IMAGE:2289660"
/clone=lib="wol_CGAP_Ut2"
/tissue_type="moderately-differentiated endometrial
adenocarcinoma, 3 pooled tumors"
/lab_host="DH10B"
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The WashU-HHMI Mouse EST Project
Unpublished (1996)
Contact: Marra M/Mouse EST Project
WashU-HHMI Mouse EST Project
WashIngton University School of Medicinep
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
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Mismatches:
Indels:
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Insert Length: 1799 Std Error: 0.00 Seq primer: -400P from Gibco High quality sequence stop: 1. Location/Qualifiers
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Matches:
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/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                       /organism="Homo sapiens"
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Fax: 314 286 1810
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                                                                                                                                                                                                                                                                                                                                              plasmid inserts
Unpublished (2000)
Unpublished (2000)
Unpublished (2000)
University of Utah Genome Center
University of Utah
Weiss
University of Utah
University Of U
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2M0113H24F Mouse 10kb plasmid UUGC1M library Mus musculus genomic close UUGC2M0113H24 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Muses 1 to 35)

1 (bases 1 to 35)

1 Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,

1 Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse whole genome scaffolding with paired end reads from 10kb
                  /clone_lib="Soares_mammary_gland_NbMMG"
/sex="male"
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7
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Mismatches:
Indels:
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                                                                                                 /tissue_type="mammary gland"
/dev_stage="4 weeks"
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t
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Seq primer: CGTTGTAAAACGACGGCCAGT
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/db_xref="taxon:10090"
/clone="UUGC2M0113H24"
/clone="IMAGE:1399836"
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 g
                                                                                                                                                                         /lab_host="DH10B"
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AZ832681.1 GI:13002685
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38.00
88.89%
77.78%
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         house mouse.
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agances gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BE395744 36 bp mRNA linear EST 21-JUL-2000 601311259F1 NIH_MGC_44 Homo sapiens cDNA clone IMAGE:3632655 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                               /lab_host="E. Coli strain XLIO-Gold, Tl-resistant, F-" /note="Weetcr: PWA2Ivy Purified genomic DNA from M. musculus C57BL/67 (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/ILNL at:
http://image.llnl.gov.f column: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 (bases 1 to 36)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
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/db_xref="taxon:9606"
/clone="IMAGE:3632655"
/clone=lib="NHH_MGC_44"
/tissue_type="endometrium, adenocarcinoma cell line"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDNA Library Preparation: Ling Hong/Rubin Laboratory CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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Matches:
Conservative:
Mismatches:
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Location/Qualifiers
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38.00
77.78%
66.67%
3.44%
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BE395744
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Best Local Similarity:
Query Match:
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JOURNAL
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA
/lab_host="DH10B (phage-resistant)"
/note="Organ: uterus; Vector: pOTB7; Site_1: XhoI; Site_2:
EGORI; CDNA made by Oligo-dr priming. Directionally
Elong into EcoRI/XhoI sites using the following 5'
adaptor: GGCACGAG(6). Library constructed by Ling Hong
in the laboratory of Gerald M. Rubin (University of
California, Berkeley) using ZAP-cDNA synthesis kit
(Stratagene) and Superscript II RT (Life Technologies)."
9 c 10 g 10 t
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., islam, H., Longarers, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly im., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AZS80135 37 bp DNA linear GSS 13-DEC-200
1M0368E16F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UGC1M036BE16"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                   Conservative:
Mismatches:
Indels:
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Insert Length: 10000 Std Error: 0.00
Plate: 0368 row: E column: 16
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AZ580135.1 GI:11694564
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38.00
83.33%
50.00%
3.44%
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Fax: 801 585 7177
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Query Match:
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84112,
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ORIGIN
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VERSION

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Score:

TITLE

COMMENT

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was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gil4732114)gblAr129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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                                                                                                                                                                                                                                                                            Edaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected for ampicillin resistance. To 20 g 3 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: cgapbs-remail.nih.gov

Tyssue Procurement: Miklos Palkovits, M.D., Ph.D.

CDNs Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
Toshiyuki and Piero Carninci (RIKEN)

CDNs Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
Sukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostoml,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
1 (bases 1 to 40)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NIH-WGC http://mgc.ncl.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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Mismatches:
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High quality sequence stop: 40.
Location/Qualifiers
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/tissue_type="hypothalamus"
/lab_host="DH108"
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/db_xref="taxon:9606"
/clone="IMAGE:4796398"
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Best Local Similarity:
Query Match:
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.05 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were adaptored but the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT 84112, USA
1821: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AZ780387 29 bp DNA linear GSS 16-FEB-2001 2M0017N12R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0017N12 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 29)
constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
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/clone="UUGC2MO017N12"
/clone_lib="Mouse 10kb plasmid UUGClM library"
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Mismatches:
Indels:
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Insert Length: 10000 Std Error: 0.00
Plate: 0017 row: N column: 12
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                   Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                       75 ProAlaAlaProGlyAlaAlaAlaGlyProAla 85 | 11 | 11 | 11 | 11 | 11 | 11 | 13 CCGAAGGCGACGGGGACCGCT 1
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                                                                                                                                                                                                                                                                              Gaps:
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Query Match:
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AUTHORS
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TITLE

COMMENT

FEATURES

AZ780387

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LOCUS

KEYWORDS

SOURCE

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of pwD42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and solected for ampicillin resistance."
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Rellly.M., Rose,M., Rose,M., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AZ784258 33 bp DNA linear GSS 16-FEB-200
2M0026G22R Mouse 10kb plasmid UUGCIM library Mus musculus genomic
clone UUGC2M0026G22 R, DNA sequence.
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Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Mismatches:
Indels:
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Matches:
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Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends stop: 33.
High quality sequence stop: 33.
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0026G22"
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Insert Length: 10000 Std Erro
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GSS.
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85.718
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Tel: 801 585 5606
Fax: 801 585 7177
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Best Local Similarity:
Query Match:
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TITLE
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polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gelectrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114 gb)RR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for amplcillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima Bonaldo
, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortlum/LLNL at:
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//note="vector: pT713D-Pac (Pharmacia) with a modified
//note="vector: pT713D-Pac (Pharmacia) with a modified
//note="vector: pT713D-Pac (Pharmacia) with a modified
//note="vector: ptyle="vector: ptyle: strand cDNA was prepared from mRNA obtained from Clontech Laboratories"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A1217993 34 bp mRNA linear EST 10-NOV-1998 qf552f10.x1 Soares_testis_NHT Homo sapiens CDNA clone IMAGE:1755698 3' similar to SW:CA11_MOUSE P11087 PROCOLLAGEN ALPHA 1(1) CHAIN PRECURSOR. ;contains MER22.b3 MSR1 MSR1 repetitive element ;; mRNA
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Mismatches:
Indels:
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/sex="male"
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Matches:
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/organism="Homo sapiens"
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100.00%
83.33%
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Best Local Similarity:
Duery Match:
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AI217993
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ORIGIN
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TITLE
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KEYWORDS
SOURCE
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gv92e02.xl NCI_CGAP_Ut2 Homo sapiens CDNA clone IMAGE:1989050 3' similar to SW:CAll_RAT P02454 COLLAGEN ALPHA 1(I) CHAIN ;contains element MER22 repetitive element ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Finsue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
Emmert-Buck, M.D., Ph.D.
Emmert-Buck, M.D., Ph.D.
Emmert-Buck, M.D., Ph.D.
End Library Arrayed by: Greg Lennon, Ph.D.
ENA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NoT-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LINL at:
www-bio.lini.gov/Dbrp/finage/image.html
Weg primer: -40UP from Gibco
High quality sequence stop: 1.
Location/Qualifiers
                                            Double-Stranded CDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT/T3 vector. Library went through one round of normalization to Cot5, and was constructed by Bento Soares and M. Fatima Bonaldo. "
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. I Chases I to 34.
NCI-CGAP http://www.ncbi.nlm.nlh.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
/clone_lib="NCI_CGAP_Ut2"
/tissue_type="moderately-differentiated endometrial
adenocarcinoma, 3 pooled tumors"
/lab_host="DH10B"
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/db_xref="taxon:9606"
/clone="IMAGE:1989050"
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Unpublished (1997)
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Query Match:
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Pred. No.:
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/note="Organ: pooled: Vector: pT7T3D-Pac (Pharmacia) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; Equal amounts of plasmid DNA from five normalized libraries were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from pools of 5,000 clones made from the same 5 libraries. The pools consisted of the following libraries and cloneIDs: Soares NbHSF pool 1: 309384-310919, 3232089-325895 Soares NDEAP pool 1: 145032-147335, 147720-148103, 148872-149255, 15002 - 150407, 151176-15237 Soares NbEHR-9W pool 1: 758280-766583, 772104-774407 Soares NbHPA pool 1: 723720-726407, 739080-740999 Subtraction by Bento Soares and M. Fatima Bonaldo."
                                                                                                                                  wg68c11.x1 Soares_NSF_F8_9W_OT_PA_P_S1 Homo sapiens cDNA clone
IMAGE:2370260 3' similar to TR:Q61888 Q61888 PROLINE RICH PROTEIN.
:Contains TAR1.t3 MSR1 repetitive element ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 37)

NCI-CGAP http://www.ncb1.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
This clone is available royalty-free through LLNL; contact the
INAGE Consortium (info@image.llnl.gov) for further information.
Seq primer: -40UP from Gibco
High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /db_xref="taxon:9606"
/clone="IMAGE:2370260"
/clone_lib="Soares_NSF_F8_9W_OT_PA_P_S1"
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          32 CCCGGGGTTTTTCCCCCCCCCCGGGG
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AI015468
                                                          RESULT 24
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LOCUS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.inh.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
NA Sequencing by: Washington University Genome Sequencing Center
found through the I.M.A.G.E. Consortium/LLNL at:
WWW-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                                                               A1612902

34 bp mRNA linear EST 07-MAR-2000 ty3303.x1 NCI_CGAP_Ut2 Home sapiens cDNA clone IMAGE:2280869 3's similar to TR:000599 00010; contains element MER22

repetitive element; mRNA sequence.
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.85 kb. Life Technologies catalog #:
11539-012"
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/clone_lib="NCI_CGAP_Ut2"
/risus_type="moderately-differentiated endometrial adenocarcinoma, 3 pooled tumors"
/lab_host="DH108"
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                                                                                                                          US-09-375-514-22 (1-205) x AI362411 (1-34)
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AI612902.1 GI:4622069
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Percent Similarity:
Best Local Similarity:
Query Match:
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AI015468 40 bp mRNA linear EST 21-DEC-1998 ov54ell.s1 Soares_testis_NHT Homo sapiens cDNA clone IMAGE:1641164 3' similar to SW:H2AL_HUMAN P98176 HYPOTHETICAL 12.7 KD HISTONE H2A RELATED PROTEIN. ;, mRNA sequence.
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10 g
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46 ProGlyIlePheSerSerGlnProGly 54

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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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AZ982483
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                                                                                                                                                                                                                                                             CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima Bonaldo
                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 40)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NAtional Cancer Institute, Cancer Genome Anatomy Project (CGAP), Thuor Gene Index Unpublished (1997)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                         1. .40
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1641164"
/clone_lib="Soares_testis_NHT"
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                     AI015468.1 GI:3229804
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Tunor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-GGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-blo.llnl.gov/bbrp/image/image.html
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2M0263K16F Mouse 10kb plasmid UUGC2M library Mus musculus genomic clone UUGC2M0263K16 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note="Organ: ovary; Vector: pCMV-SPORT6; Site_1: Sall; Site_2: Not1; Cloned unidirectionally. Primer: Oligo dT. Average insert size_1:35 kb. Tumor types include: mixed Mullerian tumor, papillary serous, clear cell, spindle cell. All are primary tumors, metastasis positive. Life technologies catalog #: 1534-013" others
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1 (bases 1 to 40)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="NCI_CGAP_OV23"
/tissue_type="tumor, 5 pooled (see description)"
/lab_host="DH10B"
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Mismatches:
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Seq primer: -400P from Gibco
High quality sequence stop: 1.
Location/Qualifiers
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/organism="Homo sapiens"
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University of Utah Genome Center
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COMMENT
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                                    E., SLC, UT
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                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="E. coli strain XLIO-Gold, Tl-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (female) was obtained from the Jackson
Laboratory Mouse DNA Resource
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (Dases 1 to 27)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Other_ESTs: 2821187.3prime
                                 S. 2030
                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Mouse 10kb plasmid UUGC2M library"
/sex="Female"
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20
84112, USA
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Mismatches:
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                                                                                                                                                    Plate: 0263 row: K column: 16
Seg primer: CGTGTAAAACGACGCCAGT
Class: plasmid ends
High quality sequence stop: 24.
Location/Qualifiers
                                                                                                                                                                                                                                                                                         /organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:10090"
/clone="UUGC2M0263K16"
                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0263 row: K column: 16
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85.71%
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3.26%
                                                                   Tel: 801 585 5606
Fax: 801 585 7177
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Email: cgapbs-remail.nih.gov
Tissue Procurement: DCTUD/DTP cDNA Library Preparation: Ling
Hong/Rubin Laboratory cDNA Library Arrayed by: The I.M.A.G.E.
Consortium (LLNL) DNA Sequencing by: Berkeley MGC Sequencing
Project Clone distribution: MGC clone distribution information can
be found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html Base Calling / Quality
Scores: PHRED from University of Washingtion Genome Center.
Trimming: cross_match from University of Washingtion Genome Center
PHRAP suite. Poly-T Identification: patwatch.pl from Berkeley
Drosophila Genome Project. University of Washingtion Genome Center:
http://www.genome.washington.edu Low Quality Sequence: 9 contiguous
PHRED high quality bases following vector sequence. Very Low
following vector sequence.
Plate: LLCMG row: C column: 12
High quality Sequence stop: 9.
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: capbs-remain.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael
Emmert-Buck, M.D., Ph.D., Michael
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Contact: Robert Strausberg,
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www-bio.llnl.gov/bbrp/image/image.html
cDNA Library Preparation: M. Bento Soares, Ph.D. CDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Brail: cgapbs-remail.nih.gov
Tissue Procurement: Christopher A. Moskaluk, M.D., Ph.D., Michael
Emmert-Buck, M.D., Ph.D.
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                             /organism="Homo sapiens"
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Matches:
Conservative:
Mismatches:
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weboco6.x1 NCI_CGAP_Pan1 Homo sapiens cDNA clone IMAGE:2324938 34 similar to SW.PSF_HUMAN P23246 PTB-ASSOCIATED SPLICING FACTOR A.contains MSR1.t3 MSR1 repetitive element; mRNA sequence.
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Email: cgapbs-rémail.nih.gov
Life Technologies catalog #: 11548-013
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
                                                                                                                                                                                                                                                                                      /note—"Vector: pr7713D-Pac (Pharmacia) with a modified polylinker; 1st strand cDNA was prepared from 3 pooled germ cell tumors, and was then primed with a Not I oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pr773 vector. Library is not normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. "
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                             /organism="Homo sapiens"
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/tissue_type="pooled germ cell tumors"
/lab_host="DH108"
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/db_xref="taxon:9606"
/clone="TMAGE:324938"
/clone=lib="NCI_CGAP_Pan1"
/tissue_type="adenocarcinoma"
/lab_host="DH108"
Trace considered overall poor quality Seg primer: -40ml3 fwd. ET from Amersham High quality sequence stop: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative:
Mismatches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length:
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High quality sequence stop: 1.
Location/Qualifiers
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                                                                                        Location/Qualifiers
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3.26%
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Unpublished (1997)
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Best Local Similarity:
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us-09-375-514-22.rst

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/note-*Organ: pooled pancreas and spleen; Vector:

pcww-sports; Site_1: Not!; Site_2: EcorV (destroyed); RNA
pourve anonymous pool of spleen and pancreas from 28 yo
male. Library is oligo-dT primed and directionally cloned
(EcorV site is destroyed upon cloning). Average insert
size 1.5 kb, insert size range 1-2.5 kb. Library is
normalized and enriched for full-length clones and was
constructed by C. Gruber (Invitrogen). Research Genetics
tracking code 025. Note: this is a NIH_MGC Library."
                                                                                                                                                                                                                                                            BI832949 32 bp mRNA linear EST 04-OCT-2001
603090781F1 NIH_MGC_120 Homo sapiens cDNA clone IMAGE:5229596 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EST 04-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. I (bases 1 to 32)
11H-MG http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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Matches:
Conservative:
Mismatches:
        Mismatches:
Indels:
Gaps:
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plate: LiAM11578 row: a column: 21
High quality sequence stop: 32.
Location/Qualifiers
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/organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /clone="IMAGE:5229596"
/clone_lib="NIH_MGC_120"
/lab_host="DH10B"
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                                                                                                                                   84 ProAlaLeuSerProValProPro 91
                                                                                                                                                         30 CCGCCTCTGTCTCCGCTGCCTCCA 7
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36.00
87.50%
75.00%
        Best Local Similarity: 75.00%
                              3.26%
                                                                                                                                                                                                                                                                                                        mRNA sequence.
BI832949
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                              Query Match:
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BI832949/c
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                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM
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KEYWORDS
SOURCE
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pcW-SPORT6: Site_1: NotI; Site_2: EcoRV (destroyed); RNA
source anonymous pool of 3 colons, age 26 yo male, 49 yo
female, 71 yo male colon; 46 yo male kidney, and pool of 2
stomachs, 62 yo male and 70 yo female. Library is
oligo-dT primed and directionally cloned (EcoRV site is
destroyed upon cloning). Average insert size 1.4 kb,
insert size range 1.3 kb. Library is normalized and
enriched for full-length clones and was constructed by C.
Gruber (Invitrogen). Research Genetics tracking code
023. Note: this is a NIH_MGC Library."
/note="Organ: pancreas; Vector: pCMV-SPORT6; Site_1: Sal1; Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT. Average insert size 1.72 kb. Life Technologies catalog #: 11548-013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BI762555 32 bp mRNA linear EST 25-SEP-2001
603048685F1 NIH_MGC_116 Homo sapiens cDNA clone IMAGE:5188724 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. I Chases 1 to 32)

NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)

Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-remail.nih.gov
Tissue Procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
Phttp://image.llnl.gov
Plate: LLAMIATI row: j column: 21
High quality sequence stop: 32.
Location/Qualiflers
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Matches:
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:5188724"
/clone_lib="NHH_MGC_l16"
/lab_host="DH10B"
                                                                                                                                                                         Length:
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BI762555.1 GI:15754133
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DB:
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BI762555/c
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Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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36.00
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Fax: 801 585 7177
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84112, USA
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AZ320746/c
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KEYWORDS
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                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo.

E 1 (bases 1 to 32)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

Lonpublished (1999)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov
Tissue Procurement: Life Technologies, Inc.

CDNA Library Preparation: Life Technologies, Inc.

CDNA Library Preparation: Life Technologies, Inc.

CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://mage.llnl.gov

Plate: LLAMI1578 row: a column: 22
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pcMv-SpORT6; Site_1: Not!; Site_2: EcoRV (destroyed); RNA
source anonymous pool of spleen and pancreas from 28 yo
male. Library is oligo-dT primed and directionally cloned
(EcoRV site is destroyed upon cloning). Average insert
size 1.5 kb, insert size range 1-2.5 kb. Library is
normalized and enriched for full-length clones and was
constructed by C. Gruber (Invitrogen). Research Genetics
tracking code C. S. Note: this is a NIH_MGC Library."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
603090881F1 NIH_MGC_120 Homo sapiens cDNA clone IMAGE:5229597 5',
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NHI-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism-"Homo sapiens"
/db_xref="taxon:9606"
/clone-ThMAGE:5229597"
/clone_lib-"NIH_MGC_120"
/lab_host="DH10B"
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                                                           BI833158.1 GI:15944708
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BI833514
                        mRNA sequence.
                                                                                                                      Homo sapiens
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Best Local Similarity:
Query Match:
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BI833514/c
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AUTHORS
TITLE
JOURNAL
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ORIGIN
    DEFINITION
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1M0040P2IR Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0040P2I R, DNA sequence.
AA320746
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pcwV-SPORT6; Site_1: NotI; Site_2: EcoNV (destroyed); RNA
pcwV-sport6; Site_1: NotI; Site_2: EcoNV (destroyed); RNA
source anonymous pool of spleen and pancreas from 28 yo
male. Library is oligo-dT primed and directionally cloned
(EcoNv site is destroyed upon cloning). Average insert
size 1.5 kb, insert size range 1-2.5 kb. Library is
normalized and enriched for full-length clones and was
constructed by C. Gruber (Invitrogen). Research Genetics
tracking code 0.55. Note: this is a NIH_MGC Library."
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Tissue procurement: Life Technologies, Inc.
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAMIS71 row: b column: 22
High quality sequence stop: 32.
High quality sequence stop: 32.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 32)
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Mouse whole genome scaffolding with paired end reads from 10kb
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Mismatches:
Indels:
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/organism="Homo sapiens"
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/clone_lib="NIH_MGC_120"
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/clone="IMAGE:2222853"
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         http://image.llnl.gov
Plate: LLAM11425 row: p
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Unpublished (1997)
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Best Local Similarity:
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/Jac-E.

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603079396F1 NIH_MGC_119 Homo sapiens cDNA clone IMAGE:5171207 5',
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Tissue Procurement: Life Technologies, Inc.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
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NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Plate: 0040 row: P column: 21
Seq primer: CACACAGGAAACAGCTATGACC
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                                                    Class: plasmid ends
High quality sequence stop: 32.
Location/Qualifiers
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                                                                                                                                                                                                        /strain="C57BL/6J
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36.00
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DEFINITION
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AUTHORS
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/note="Organ: brain; Vector: pCMV-SPORT6; Site_1: NotI; Site_2: ECGNV (destroyed); RNA source normal medulla from anonymous male age 27. Library is oligo-dr primed and directionally cloned (ECGNV site is destroyed upon cloning). Average insert size 1.3 kb, insert size range 0.9-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 013. Note:
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34 bp mRNA linear EST 21-APR-1999
tr6lhll.x2 NCI_CGAP_Panl Homo sapiens cDNA clone IMAGE:2222853 3'
similar to TR:004154 Q04154 SALIYARY PROLINE-RICH PROTEIN RP15
PRECURSOR. ;contains element MSR1 repetitive element ;; mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Life Technologies catalog #: 11548-013
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LINL at:
www-bio.llnl.gov/bbrp/image/image.html
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 34)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length:
Matches:
Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (Invitrogen). Research Genethis is a NIH_MGC Library."
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                                                                                                                                             /clone="IMAGE:5171207"
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/tissue_type="medulla"
/lab_host="DH10B"
                                                                                           /organism="Homo sapiens"
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps:
High quality sequence stop: 33.
Location/Qualifiers
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                                                                                                                    /db_xref="taxon:9606"
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High quality sequence stop: 1.
Location/Qualifiers
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36.00
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us-09-375-514-22.rst

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175 LeuTrpMetThrGluTyrLeuAsnArgHis 184
                                                            2 CTCTGGATGACATCATTGCTCAGCTCCCAC 31
US-09-375-514-22 (1-205) x BE867678 (1-35)
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Fax: 801 585 7177
                                                                                                                                                                                                                                                                           house mouse.
                                                                                                                                                                                                                    AZ477888
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Pred. No.:
Score:
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ORIGIN
                                                                                                                                       AZ477888/C
                                                                                                                                                                            DEFINITION
                                                                                                                                                                                                                                                                                             ORGANISM
                                                                                                                     RESULT 40
                                                                                                                                                                                                                    ACCESSION
                                                                                                                                                                                                                                                                                                                                                      REFERENCE
                                                                                                                                                                                                                                                                                                                                                                        AUTHORS
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                                                                                                                                                                                                                                    VERSION
                                                                                                                                                                                                                                                                           SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lucaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo.

E 1 (bases 1 to 35)

NIH-MGC http://mgc.nci.nih.gov/.

National Institutes of Health, Mammalian Gene Collection (MGC)

National Institutes of Health, Mammalian Gene Collection (MGC)

Ontact: Robert Strausberg, Ph.D.

Email: cgapbs-rémail.nih.gov

Tissue procurement: ATCC

CDNA Library Preparation: Life Technologies, Inc.

CLONA Gistribution: MGC Clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

http://image.lln.gov

Plate: LLAM9561 row: p column: 06

High quality sequence stop: 35.

Location/Qualifiers
  /tissue_type="adenocarcinoma"
/lab_host="DH10B"
/note="Organ: pancreas; Vector: pCMV-SPORT6; Site_1: Sal1;
Site_2: Not1; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1:72 kb. Life Technologies catalog #:
11548-013"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BE867678 35 bp mRNA linear EST 20-OCT-2000 601443224F1 NIH_MGC_65 Homo sapiens cDNA clone IMAGE:3847541 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="laxon:9606"
/clone=lib="NHH MGC_65"
/tissue_type="adenocarcinoma"
/lab_host="PHIOB (phage-resistant)"
/note="Organ: colon; Vector: pCMV-SPORT6; Site_1: Not1;
/note="Organ: colon; Vector: pCMV-SPORT6; Site_1: Not1;
Site_2: Salı; cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.8 kb. Library constructed by Life
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Matches:
Conservative:
Mismatches:
Indels:
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Matches:
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BE867678
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Query Match:
DB:
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Best Local Similarity:
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ORIGIN
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JOURNAL
COMMENT
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KEYWORDS
SOURCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 39
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarcse gelectrophoresis. Vector DNA was prepared from a derivative of pWD42 (qil4732114) plant29072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for amploillin resistance."
                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 35)
SS Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
In Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
AZ477888 35 bp DNA linear GSS 04-OCT-2000 1M0297B02R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0297B02 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT 84112, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /lab_host="E. Coli strain XLI0-Gold, Tl-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UGC1M0297B02"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0297 row: B column: 02
Seg primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Laboratory Mouse DNA Resource
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5.68e+05
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qu01c09.x1 NCI_CGAP_CO14 Homo sapiens cDNA clone IMAGE:1963504 3'
sinilar to SW.EXTN DAUCA P06599 EXTENSIN PRECURSOR.; contains
A1357425
A1357425.1 GI:4109046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /tissue_trype="kidney tumor"
/lab_host="SOLR (kanamycin resistant)"
/note="Yogan: kidney; Vector: Bluescript SK-; Site_1:
BCORI; Site_2: XhOI; Cloned unidirectionally. Primer:
Oligo dr. Pooled kidney tumors. 5' adaptor sequence: 5'
GAATTCGCACCAG 3' 3' adaptor sequence: 5'
CTCGATTTTTTTTTTTTTTTT 3' Average insert size: 1.0 kb."

a 1 c 19 g 13 t
                                                                                                                                                                                                                                                                     Tunor Gene Index

Tunor Gene Index

Uppublished (1997)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

Tissue Procurement: L. Jeffrey Medeiros, M.D., Michael R.

Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: Stratagene, Inc.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
           og87g06.sl NCI_CGAP_Kid6 Homo sapiens cDNA clone IMAGE:1593370 3' similar to TR:099322 Q99322 MUCIN CORE POLYPEPTIDE, TRACHEAL AA987401
                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 37)
                                                                                                                                                                                                                                    NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           37
1
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  www-bio.llnl.gov/bbrp/image/image.html
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Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone="IMAGE:1593370"
/clone_lib="NCI_CGAP_Kid6"
/sex="mixed"
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/organism="Homo sapiens"
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                                                                                          AA987401
AA987401.1 GI:3172765
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                                                                                                                                                                   Homo sapiens
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                                      DEFINITION
   AA987401/c
                                                                                                                                                                                                                                    AUTHORS
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AI357425
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                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.

1 (bases 1 to 36)
Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,P., Zimmerman,J. and Ecker,J.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: ecker@salk.edu
This is single pass sequence recovered from the left border of
TDNA. This sequence lies within an annotated intron of At5g52330.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Arabidopsis Genome Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
   Conservative:
Mismatches:
Indels:
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Mismatches:
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                                                                                          US-09-375-514-22 (1-205) x AZ477888 (1-35)
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36.00
87.50%
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                                                                                                                                                                                                                                                                                                                                                                                      Arabidopsis thaliana
100.00%
83.33%
3.26%
                                                                                                                              55 HisThrProHisProAla 60
                                                                                                                                              34 CACACCCCCACCCAGC 17
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                                                                                                                                                                                                                                                                                                                                                                     thale cress.
 Percent Similarity:
Best Local Similarity:
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DB:
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VERSION
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BH792266
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CDNA Library Proparation: Life Technologies, Inc.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1963504"
/clone="IMAGE:1963504"
/tibsue_type="moderately-differentiated adenocarcinoma"
/tibsue_type="moderately-differentiated adenocarcinoma"
/lab_host="DH10B"
/note="Organ: colon; Vector: pcMV-SPORT6; Site_1: SalI;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.7 kb. Life Technologies catalog #:
11531-019"
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 37)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

    (bases 1 to 38)

                                                                        NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael
Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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Contact: Robert Strausberg, Ph.D.
Emall: cgapbs-remail.nih.gov
Tissue Procurement: Life Technologies, Inc.
CDM Library Preparation: Life Technologies, Inc.
CDM Library Arrayed by: The I.M.A.G.E. Consortium (LINL)
DNA Sequencing by: Incyte Genomics, Inc.
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Mismatches:
Indels:
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Location/Qualifiers
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Matches:
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                                                                                                                                                                  Contact: Robert Strausberg, Ph.D.
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BI829358
BI829358.1 GI:15940908
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63.64%
63.64%
3.26%
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Unpublished (1997)
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BI829358/c
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                                                  REFERENCE
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/clone_lib="NIH_MGC_119"
/tissue_type="medulla"
/tissue_type="medulla"
/tissue_type="medulla"
/tissue_type="medulla"
/tissue_type="medulla"
/note="Organ: brain; Vector: pCMV-SPORT6; Site_1: NotI;
Site_2: EcoRV (destroyed); RNA source normal medulla from anonymous male age 27. Library is oligo-dT primed and directionally cloned (EcoRV Site is destroyed upon cloning). Average insert size 13 kb, insert size range 0.9-3 kb. Library is normalized and enriched for full-length clones and was constructed by C. Gruber (Invitrogen). Research Genetics tracking code 013. Note: this is a NIH_MC Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AA869375 40 bp mRNA linear EST 16-MAR-1998 vq07c10.rl Barstead stromal cell line MPLRB8 Mus musculus cDNA clone IMAGE:1093554 5' similar to TR:014883 Q14883 INTESTINAL MUCIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 40)
Marra, M., Hillier, L., Allen, M., Bowles, M., Dietrich, N., Dubuque, T., Geisel, S., Kucaba, T., Lacy, M., Le, M., Martin, J., Morris, M., Schellenberg, K., Steptce, M., Tan, F., Underwood, K., Moore, B., Theising, B., Wylie, T., Lennon, G., Soares, B., Wilson, R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This clone is available royalty-free through LLNL ; contact the IMAGE Consortium (info@image.llnl.gov) for further information.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: LLAM11425 row: f column: 23 High quality sequence stop: 38.

Location/Qualifiers
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Unpublished (1996)
Contact: Marra M/Mouse EST Project
WashU-HHMI Mouse EST Project
WashU-HHMI Mouse EST Project
WashIngton University School of Medicinep
WA44 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1810
Fax: 314 286 1810
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Possible reversed clone: similarity on wrong strand
Seq primer: -20m13 rev2 ET from Amersham
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Mismatches:
Indels:
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Matches:
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                                                                                                                                                                                                       /db_xref="taxon:9606"
/clone="IMAGE:5170966"
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87.50%
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3.26%
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AA869375
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Best Local Similarity:
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GSS 02-OCT-2000
                                                   /clone_InAGR:212867"
/clone_Lib="NCI_CGAP_Utl"
/clone_Lib="NCI_CGAP_Utl"
/clone_Lib="NCI_CGAP_Utl"
/clone_Lipe="well-differentiated endometrial
adenocarcinoma, 7 pooled tumors"
/lab_host="hulbs"
/note="Organ: uterus; Vector: pCMV-SPORT6; Site_1: SalI;
Site_2: NotI; Cloned unidirectionally. Primer: Oligo dT.
Average insert size 1.75 kb. Life Technologies catalog #:
11538-014"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bukaryota, Metazoa, Chordata, Craniata, Vertebrata; Euteleostomi;
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae, Murinae; Mus.

I (bases 1 to 40)
Dunn, D., Aoyagi, A., Barber, M., Bacorn, T., Duval, B., Hamil, C.,
Islam, H., Longaree, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
'M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A2372178 40 bp DNA linear GSS 02-OCT-200 1M0123J24R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0123J24 R, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (2000)
Contact: Robert B. Weiss
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
84112, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="E. Coli strain XLIO-Gold, Tl-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/61 (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
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7
7
8
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Mismatches:
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Fax: 801 585 7177

Email: ddun@qenetics.utah.edu
Insert Length: 10000 Std Error: C
Plate: 0123 row: J column: 24
Seg primer: CACACAGGAAACAGCTATGACC
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                         /organism="Homo sapiens"
/db_xref="taxon:9606"
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/organism="Mus musculus"
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/clone="UUGC1M0123J24"
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66.67%
58.33%
3.26%
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AUTHORS
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                                                                                                                                                                             tg56182 NCI_CGAP_Utl Homo sapiens cDNA clone IMAGE:2212867 3' similar to TR:Q01943 G01943 EXTENSIN ; contains element TARI AI555182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
CLone distribution: NCI-CGAP Clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 40)
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: cgapbs-remail.nih.gov
Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael
Emmert-Buck, M.D., Ph.D.
                                                                             /db_xref="taxon:10090"
/clone="IMAGE:1093554"
/clone_llb="Barstead stromal cell line MPLRB8"
/clone_llb="barstead stromal cell line MPLRB8"
/lab_host="DH10B"
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Mismatches:
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                                                              /organism="Mus musculus"
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High quality sequence stop: 1.
Location/Qualifiers
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Seq primer: -400P from Gibco
High quality sequence stop: 1
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71.43%
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Unpublished (1997)
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Best Local Similarity:
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KEYWORDS
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                                                                                          Pred.
               (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligomucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gil4732114[gb]AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/lab_host="DH10B (phage-resistant)"
/lab_host="DH10B (phage-resistant)"
/note="Organ: colon; Vector: pCWV-SPORT6; Site_1: Not1;
Site_2: Sal1; Cloned unidirectionally. Primer: Oligo dT.
Swerage insert size 1.8 kb. Library constructed by Life
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
Vakaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Bukammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 38)
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Tissue Procurement: ATCC
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
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http://image.llnl.gov
Plate: LLAM9561 row: e column: 09
High quality sequence stop: 38.
Location/Qualifiers
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Matches:
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/clone="IMAGE:3847280"
/clone_lib="NIH_MGC_65"
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BE867536
BE867536.1 GI:10316312
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KEYWORDS
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TITLE
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Score:

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CONA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
cDNA Library Preparation: M. Bento Soares, Ph.D., M. Fatima Bonaldo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryorta: Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. I Chases 1 to 25. NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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17
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Seg primer: -40ml3 fwd, ET from Amersham
High quality sequence stop: 1.
Location/Qualifiers
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Mismatches:
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/clone_lib="Soares_testis_NHT"
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/organism="Homo sapiens"
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Unpublished (1997)
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(http://www.jax.org/resources/documents/dhares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gilg19129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chamically completent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                   AZ791470 30 bp DNA linear GSS 16-FEB-2001
2M0041H21F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0041H21 F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 30)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /lab_host="E. Coli strain XLI0-Gold, Tl-resistant, F-" /note="Vector: PW042nv; Puiffied genomic DNA from M. musculus G57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts Unpublished (2000)
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/clone_lib="Mouse 10kb plasmid UUGCIM library"
/sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0041 row: H column: 21
Seg primer: CGTTGTAAAACGACGGCCAGT
  Indels:
Gaps:
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/strain="C57BL/6J"
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University of Utah Genome Center
University of Utah
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High quality sequence stop: 30.
Location/Qualifiers
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Fax: 801 585 7177
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ORIGIN

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Mismatches:
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Matches:
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US-09-375-514-22 (1-205) x AZ791470 (1-30)

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Copyright (c) 1993 - 2003 Compugen Ltd.
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Sequence 36, Sequence 29, Sequence 24, Sequence 147, Sequence 66, Sequence 67, Requence 69, Sequence 30, Sequence 24, Sequence 18, Sequence 18,		equence 11553 equence 11554 equence 11554 Sequence 3926 Sequence 3927 Sequence 44114 Sequence 1325	nence in ence
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18     9     US-09-808-326-756     Sequence 75i       18     9     US-09-931-732-20     Sequence 20       18     9     US-10-118-918-55     Sequence 1,       18     9     US-10-112-653-1     Sequence 1,       18     9     US-10-112-653-85     Sequence 85       18     9     US-10-017-995-1     Sequence 1,       18     9     US-10-017-995-54     Sequence 1,       18     9     US-10-017-995-54     Sequence 55	18 9 US-10-017-995-91 Sequence 91 18 9 US-09-776-479-1 Sequence 1, 18 9 US-09-776-479-54 Sequence 54 18 9 US-09-776-479-55 Sequence 55	18 9 US-09-776-479-91 Sequence 91 B 9 US-10-002-884A-6 Sequence 6, 18 9 IG-10-300-884A-6 Sequence 51 B 9 IG-10-300-51	18 10 US-09-734-846-68 Sequence 6 18 10 US-09-824-46-59 Sequence 5 18 10 US-09-824-46-59 Sequence 5 18 10 US-09-824-46-59	18 10 US-03-624-408-104 Sequence 1	18 10 US-09-965-116A-77 Sequence 7 18 10 US-09-965-116A-98 Sequence 9	18 10 US-09-965-116A-99 Sequence 9 20 9 US-10-085-056-12 Sequence 12	20 9 US-09-888-326-757 Sequence 75 20 9 US-09-888-326-766 Sequence 76	20 9 US-09-932-300-72 Sequence /2 20 9 US-10-112-653-81 Sequence 81	20 9 US-10-112-653-102 Sequence 10 20 9 US-10-017-995-87 Sequence 87	20 9 US-10-017-995-109 Sequence 10 20 9 US-09-776-479-87 Sequence 87	20 9 US-09-776-479-109 Sequence 10 21 9 US-10-074-246-14 Sequence 14	21 10 US-09-1/65-081-598 Sequence 3 21 10 US-09-817-014-61 Sequence 6 24 9 US-09-840-185-3361 Sequence 33	25 9 US-09-745-317-41 Sequence 41	25 5 00 10 21 10 11 11 11 10 10 10 10 10 10 10 10 10	25 10 US-09-866-108-5162 Sequence 5	25 10 US-09-399-111 Sequence 1	26 9 US-10-006-856A-92 Sequence 92	26 9 US-10-006-818A-92 Sequence 92 26 9 US-10-015-393A-92 Sequence 92	26 9 US-09-946-374-92 Sequence 92	26 9 US-10-015-869A-92 Sequence 92	26 9 US-1U-0U6-116A-92 Sequence 92 26 9 US-1O-006-117A-92 Sequence 92	26 9 US-10-013-913A-92 Sequence 92	26 9 US-10-007-194A-92 Sequence 92	26 9 US-10-013-430A-92 Sequence 92 26 9 US-10-011-671A-92 Sequence 92	26 9 US-10-012-755A-92 Sequence 92 28 9 US-09-345-373-92 Sequence 92	28 9 US-09-345-373-114 Sequence 11	28 10 US-09-853-666-13 Sequence 1 29 9 US-09-991-936-51 Sequence 51	29 10 US-09-817-387-12 Sequence	30 9 US-09-953-052-50 Sequence 50 30 9 US-10-196-793A-40 Sequence 40	30 10 US-09-804-682-80 Sequence	31 9 US-10-158-895-29 Sequence 29 31 9 US-09-939-833-1 Sequence 1,	31 9 US-09-912-263-108 Sequence 10	31 10 0S-10-23-0/4-35 31 10 0S-09-9801-31 32-980-9801-274-112 Sequence 3-31 10 0S-09-9801-31 Sequence
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9 18 9 US-09-988-326-756 Sequence 75i 9 18 9 US-09-931-732-20 Sequence 20 9 18 9 US-09-918-55 Sequence 20 9 18 9 US-10-112-653-1 Sequence 1, 9 18 9 US-10-112-653-1 Sequence 1, 9 18 9 US-10-017-995-1 Sequence 85 9 18 9 US-10-017-995-54 Sequence 54, 9 18 9 US-10-017-995-54 Sequence 54	2.9 18 9 US-10-017-995-91 Sequence 91 2.9 18 9 US-09-776-479-1 Sequence 1, 2.9 18 9 US-09-776-479-54 Sequence 54 2.9 18 9 US-09-776-479-55 Sequence 55 2.9 18 9 US-09-776-479-55	2.9 18 9 US-09-776-479-91 Sequence 91 2.9 18 9 US-10-012-848.4-6 Sequence 6, 9 18 9 IS-10-301-347-51 Sequence 51 9 18 9 IS-10-301-347-51 Sequence 51	2.9 18 10 US-09-734-846-68 Sequence 6 2.9 18 10 US-09-824-468-59 Sequence 5	2.9 18 10 US-09-969-173-4373 Sequence 7 2.9 18 10 US-09-965-116A-7 Sequence 7	2.9 18 10 US-09-965-116A-77 Sequence 7 2.9 18 10 US-09-965-116A-98 Sequence 9	2.9 18 10 US-09-965-116A-99 Sequence 9 2.9 20 9 US-10-085-056-12 Sequence 12	2.9 20 9 US-09-888-326-757 Sequence 75 2.9 20 9 US-09-888-326-766 Sequence 76	2.9 20 9 US-09-932-300-72 Sequence /2 2.9 20 9 US-10-112-653-81 Sequence 81	2.9 20 9 US-10-112-653-102 Sequence 10 2.9 20 9 US-10-017-995-87 Sequence 87	2.9 20 9 US-10-017-995-109 Sequence 10 2.9 20 9 US-09-776-479-87 Sequence 87	2.9 20 9 US-09-776-479-109 Sequence 10 2.9 21 9 US-10-074-246-14 Sequence 14	2.9 21 10 US-09-765-081-398 sequence 3 2.9 21 10 US-09-817-0114-61 Sequence 6 2 0 24 0 HS-09-040-185-3351 Sequence 33	2.9 25 9 US-09-745-317-41 Sequence 41	2.9 25 10 0S-09-866-108-4386 Sequence 4 2 2 25 10 0S-09-866-108-4386 Sequence 4 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	2.9 25 10 US-09-866-108-5162 Sequence 5	2.9 25 10 US-09-398-399-11 Sequence 1	2.9 26 9 0S-10-006-856A-92 Sequence 92	2.9 26 9 US-10-006-818A-92 Sequence 92 2.9 26 9 US-10-015-393A-92 Sequence 92	2.9 26 9 US-09-946-374-92 Sequence 92	2.9 26 9 US-10-015-869A-92 Sequence 92	2.9 26 9 US-10-006-116A-92 sequence 92 2.9 26 9 US-10-006-117A-92 sequence 92	2.9 26 9 US-10-013-913A-92 Sequence 92	2.9 26 US-10-07-194A-92 Sequence 92	2.9 26 9 US-10-013-450A-92 Sequence 92 2.9 26 9 US-10-011-671A-92 Sequence 92	2.9 26 9 US-10-012-755A-92 Sequence 92 2.9 28 9 US-09-345-373-92 Sequence 92	2.9 28 9 US-09-345-373-114 Sequence 11	2.9 28 10 US-09-853-666-13 sequence 1 2.9 29 9 US-09-991-936-51 Sequence 51	2.9 29 10 US-09-817-387-12 Sequence 1	2.9 30 9 US-09-953-052-50 Sequence of 2.9 30 9 US-10-196-793A-40 Sequence 40	2.9 30 10 US-09-804-682-80 Sequence 8	2.9 31 9 US-10-158-895-29 Sequence 29 2.9 31 9 US-09-939-833-1 Sequence 1,	2.9 31 9 US-09-912-263-108 Sequence 10	2.9 31 9 US-10-235-0/4-25 Sequence 3- 2.9 31 10 US-09-801-274-112 Sequence 1 2.0 31 10 US-08-801-274-313 Seminore 6

Sequence 5, Appli Sequence 3, Appli Sequence 37, Appl		10,4	21,	21,	302,	Sequence 17	<b>س</b> د		_` `	Sequence 87, Appl	5	100	~ «	,	Sequence 51, Appl	Sequence 5023, Ap	ò	Sequence 31, Appl	382	9,	18,	13,	47	Sequence 47, Appl Sequence 47, Appl		5,6	30	320	Sequence 30, Appl	Sequence 11, Appli	14,	Sequence 141, App	Sequence 61, Appl	6									N THE	CARDIOVASCULAR DISEASE	
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C 442 444	445 446 447	c 448 449	450	1.4	c 453	4 -	c 457	458	4 4	c 461	463	4	466	46	469	c 470	471	473	47	C 4/5	47	c 478 479	480	481	483	c 485	4.4	4	C 489	4.	4 4 9 2 2 9 3	494	4 4	C 497	4 4	. r			RESULT 1	US-09-970	; Sequence 11, Application of Patent No. US20020170077A1	GENER		•••	
2697, Ap 2698, Ap 2699, Ap 855	2863	3509	9820	4533	3928,	ce 3929, Ap	4388,	4389,	5606,	ce 5607, Ap	5609,	5610,	11102,	13837,	13839,	13840,	13841,	139	927	928	928	ce 930, App	931	17,	<b>A</b>	~~	4	٦,	8, Appli 8, Appli	24	11, Appi 8, Appli	e 3, Appli	5449	13, Appl	680	1046,	1340, Ap	17, Appl	54, Appl 60, Appl	54, Appl	35,	89, Appl 15, Appl	7	17	83,
Sequence					uenbes Sednen	Sequen	Sequen	Sequen	Sequen	Sequen	Sequen	Sequen	Sequen	Sequen	Sednen	Sednen	Sequen	Sednen	Sequen	nenbes Sednen	Sequen	gednen	Sequen	Sequen	Sequence	1 01	Sequence	Sednence	Sequence	Sequence	Sednence	Seguenc	Sequence	Seguenc	Sednenc	Sequence	Sequenc	Sequence	Sequence	Sequence	Sequence	Sequence	ໝະ	Sequence	
9 US-10-060-756A-2697 9 US-10-060-756A-2698 9 US-10-060-756A-2699 0S-10-215-112-855	US-10-215-112-2863 US-10-215-112-3003	US-10-215-112-3509 US-10-215-112-3635	US-10-215-112-9820 US-10-215-112-9932	US-09-940-185-4533	US-09-866-108-3928	US-09-866-108-3929 US-09-866-108-3930	US-09-866-108-4388	US-09-866-108-4389 US-09-866-108-4390	US-09-866-108-5606	US-09-866-108-5608 US-09-866-108-5608	us-09-866-108-5609	US-09-866-108-5610 US-09-866-108-11101	US-09-866-108-11102	US-09-866-108-13837 US-09-866-108-13838	US-09-866-108-13839	US-09-866-108-13840	US-09-866-108-13841 US-09-866-108-13917	US-09-866-108-13918	US-09-827-998-927 US-09-827-998-927	US-09-827-998-928	US-09-827-998-928	US-09-827-998-930	US-09-827-998-931	US-09-918-702-17	US-10-085-476-3 US-10-201-964-5	US-09-902-460-17	US-10-231-426-4 US-09-739-451-14	US-09-983-657-1	US-09-877-336-8	US-09-952-060-24	US-10-170-997-8	US-09-952-677-3	US-09-780-533A-5449	US-09-766-396-13	US-09-801-274-680	US-09-801-274-1046	US-10-062-375-13	US-09-854-356-17	US-09-957-483-60	US-10-008-721-54 US-10-112-488-56	US-09-814-786-35	US-10-027-961A-15	US-10-022-832-57 US-10-196-730-9	US-10-252-408-11	68-626-479-60-60
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369 370 371 372	c 373			c 379			384	386		389			393	395	396	398	c 399		c 402		c 404	c 406	c 407	c 409	411	c 412	414	415			2 420		c 423	424	c 426	427 c 428		430	c 432		c 435				:

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Length:
Matches:
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APPLICATION NUMBER: US 08/599,654
FILING DATE: 09-FEB-1996
APPLICATION NUMBER: US 08/485,573
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APPLICATION NUMBER: US 08/386,844
FILING DATE: 10-FEB-1995
ATTORNEY/AGENT INFORMATION:
NAME: CORUZZI, LAURA A
                                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL: NO SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                           TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 11-Aug-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence II, Application US/09371900
Patent No. US20020137700A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         140 AspGlyValAsnTrpGlyArg 146
                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
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44.00
100.008
100.008
3.998
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STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 3
US-09-371-900-11/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: FALB, DEAN A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/970,820
FILING DATE: 05-Oct-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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Mismatches:
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ADDRESSEE: PENNIE & EDMONDS
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: PENNIE & EDMONDS
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                         NAME: Coruzzi, Laura A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7853-032
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9990
TELEFAX: (212) 869-8864
                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/386,844
FILING DATE: 10-FEB-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels:
                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL: NO
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 11, Application US/09986718 Patent No. US20020178458A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               140 AspGlyValAsnTrpGlyArg 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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                                                       New York
                                                                                           ZIP: 10036-2711
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STATE: New York
                                     CITY: New York
                                                                         COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-986-718-11/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-970-820-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 2
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMFUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATEMIT Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/371,900
Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-375-514-22 (1-205) x US-09-986-718-11 (1-24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: PENNIE & EDMONDS LLP
STREET: 1155 Avenue of the Americas
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Sequence 3, Application US/09925674A

Sequence 3, Application US/09925674A

Patent No. US20020119943A1

GENERAL INFORMATION:

APPLICANT: AMRAD Operations Pty Ltd

TITLE OF INVENTION: A NOVEL MAMMALIAN GENE, bcl-w, BELONGS TO THE bcl-Z

TITLE OF INVENTION: A MOVEL MAMMALIAN GENE, bcl-w, BELONGS TO THE bcl-Z

TITLE OF INVENTION: A MOVEL MAMMALIAN GENE, bcl-w, BELONGS TO THE bcl-Z

TITLE OF INVENTION: A NOVEL MAMMALIAN GENE, bcl-w, BELONGS TO THE bcl-Z

TITLE OF INVENTION: A NOVEL MAMMALIAN GENE, bcl-w, BELONG TO THE bcl-Z

TITLE OF INVENTION UNMBER: US/09/925,674

PRIOR FILING DATE: 2001-08-09

PRIOR FILING DATE: 1996-03-27

NUMBER OF SEQ ID NOS: 9

SOFTWARE: PATENTIN VEF. 2.1

SEQ ID NO 3

LENGHAM

LENGHAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Cargill, Michele
APPLICANT: Ireland, James S.
APPLICANT: Lander. Erics S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2009-001
CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILING DATE: 2001-03-07
PRIOR FILING DATE: 2000-03-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         00700
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 60/206,129
PRIOR FILING DATE: 2000-05-22
NUMBER OF SEQ ID NOS: 1802
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 465
LENGTH: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-801-274-465; Sequence 465, Application US/09801274; Patent No. US20020032319A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             188 TrpileGlnAspAsnGlyGlyTrp 195
     28 AACCAGCATTTACATCCTTGGATCCGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION: 14
OTHER INFORMATION: n is inosine
NAME/KEY: modified_base
LOCATION: 17
OTHER INFORMATION: n is inosine
NAME/KEY: modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTHER INFORMATION: n is inosine US-09-925-674A-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.5e+03
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75.00%
75.00%
3.71%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
US-09-801-274-465
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Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Mouse
                                                    RESULT 5
US-09-925-674A-3/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Nucleotide sequence encoding the peptide derived from a partial H OTHER INFORMATION: unan c-Myc-tag fused to the N-terminus of Zucchini Yellow Mosaic OTHER INFORMATION: Virus (AGII) coat protein gene of which the N' terminal 43 amino OTHER INFORMATION: acids were deleted
                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: other nucleic acid.
DESCRIPTION: /desc = "synthetic oligonucleotide"
HYPOTHERICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0017
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-375-514-22 (1-205) x US-09-371-900-11 (1-24)
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Matches:
REGISTRATION NUMBER: 30,742
REFRENCE/DOCKET NUMBER: 7853-104
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864
TELEFX: 66141 PENISE
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels:
                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps:
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             140 AspGlyValAsnTrpGlyArg 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23 GATGGGGTGAACTGGGGGAGG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 33
SOFWARE: Patentin version 3.1
SEQ ID NO 18
LENGTH: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
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44.00
100.00%
100.00%
3.99%
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42.00
88.89%
66.67%
3.80%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Alignment Scores:
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Sequence 372, Application US/09801274
GENERAL INFORMATION:
APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2009-001
CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILING DATE: 2001-03-07
PRIOR PLICATION NUMBER: US 60/187,510
PRIOR PLICATION NUMBER: US 60/206,129
PRIOR PLILING DATE: 2000-05-22
NUMBER OF SEQ ID NOS: 1802
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21, Application US/09931732
Publication No. US20030045488A1
GENERAL INFORMATION:
APPLICANT: Brown, Bob D.
APPLICANT: Riley, Timothy A.
TITLE OF INVENTION: ANTIENSE OLIGONUCLECTIDES COMPRISING
TITLE OF INVENTION: UNIVERSAL AND/OR DEGENERATE BASES
FILE REFERENCE: OASBIO.001C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1110881
                00000
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Mismatches:
Indels:
                                                Conservative:
Mismatches:
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                                                                                                                                         US-09-375-514-22 (1-205) x US-09-801-274-465 (1-31)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/931,732
CURRENT FILING DATE: 2001-08-16
PRIOR APPLICATION NUMBER: PCT/US00/09293
PRIOR FILING DATE: 2000-04-07
PRIOR APPLICATION NUMBER: US 60/128,377
PRIOR FILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 30
SOFTWARE: FastSEQ for Windows Version 4.0
SSOFTWARE: FastSEQ for Windows Version 21
LENGTH: 22
                                   Matches:
                                                                                       Indels:
                Length:
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38.50
80.00%
80.00%
3.49%
                5.61e+03
                                 39.00
77.78%
55.56%
3.53%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Homo sapiens
US-09-801-274-372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Percent Similarity:
Best Local Similarity:
Query Match:
                                          Percent Similarity:
Best Local Similarity:
Query Match:
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Alignment Scores:
Pred. No.:
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                                                                                                                                                                                                                                                 RESULT 7
                                     Score:
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US-09-984-842-2/C

| Sequence 2, Application US/09984842|
| Sequence 2, Application US/09984842|
| Sequence 2, Application No. US20030077592a1|
| GENERAL INPORMATION:
| APPLICANT: MORTEN, JOHN EDWARD NORRIS
| TITLE OF INVENTION: DIAGNOSTIC USE
| TITLE OF INVENTION: DIAGNOSTIC USE
| TITLE OF INVENTION: UNMBER: US/09/984,842|
| CURRENT FILING DATE: 2001-10-31|
| PRIOR PPLICATION NUMBER: 09/485,636|
| PRIOR APPLICATION NUMBER: 09/485,636|
| PRIOR APPLICATION NUMBER: 097-08-22|
| PRIOR FILING DATE: 1997-08-22|
| PRIOR FILING DATE: 1998-08-18|
| NUMBER OF SEQ ID NOS: 26|
| SOFTWARE: Patentin Ver: 2.1|
| SEQ ID NO 2|
| LENGTH: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 943, Application US/09801274
Patent No. US20020032319A1
GENERAL INFORMATION:
APPLICANT: Cargill, Michele
APPLICANT: Lander, Ente S.
APPLICANT: Lander, Ente S.
APPLICANT: Lander, Ente S.
APPLICANT: LANGER: ENTER S.
APPLICANT: WINNERTION: HUMAN SINGLE NUCLECTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2009-001
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US 60/187,510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              00100
                    ; FEATURE:
; OTHER INFORMATION: Synthetic oligonucleotide primers
US-09-931-732-21
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Mismatches:
Indels:
                                                                                                            Length:
Matches:
Conservative:
                                                                                                                                                                    Mismatches:
                                                                                                                                                                                                                                              US-09-375-514-22 (1-205) x US-09-931-732-21 (1-22)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-375-514-22 (1-205) x US-09-984-842-2 (1-30)
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Matches:
                                                                                                                                                                                        Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29 GCCCGCCTGGAGCGCCCCCGGCC
                                                                                                                                                                                                                                                                                                      2 GIGCCACCIGIGGICCACCIG 22
                                                                                                                                                                                                                                                                                  89 ValProProValValHisLeu 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6.83e+03
38.00
87.50%
87.50%
ORGANISM: Artificial Sequence
                                                                                                          4.74e+03
38.00
100.008
100.008
                                                                                                                                          Percent Similarity:
Best Local Similarity:
Query Match:
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Query Match:
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US-09-801-274-943/C
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                                                                                                Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No.:
                                                                                                                Pred. No.:
                                                                                                                                      Score:
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Sequence 370, Application US/10125635A
Fublication No. US20030039635A1
GENERAL INFORMATION:
APPLICANT: Galger, Alexander
APPLICANT: Galger, Barick
APPLICANT: Garter, Darrick
APPLICANT: McNeill, Patricia D.
APPLICANT: Stiberland, R. Alec
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR WTI
TITLE OF INVENTION: COMPOSITION AND METHODS
TITLE REFERENCE: 210121.4657
CURRENT APPLICATION NUMBER: US/10/125,635A
CURRENT FILING DATE: 2002-07-19
NUMBER OF SEQ ID NOS: 461
SEQ ID NO 370
LENGTH: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 38
7
0
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Matches:
Conservative:
Mismatches:
Indels:
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                  US-09-375-514-22 (1-205) x US-09-801-274-943 (1-31)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            36 GlyAlaAlaProProGlyAlaAlaProAlaPro 46
PRIOR FILING DATE: 2000-03-07
PRIOR APPLICATION NUMBER: US 60/206,129
PRIOR FILING DATE: 2000-05-22
NUMBER OF SEQ ID NOS: 1802
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 943
LENGTH: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 35 GGTCCAGCACCTCGCCCGCAACGCCCCCACCG 3
                                                                                                                                                                                                                                                                                                                    Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 370, Application US/09938864; Publication No. US20030072767A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Gaiger, Alexander
APPLICANT: McNeill, Patricia D.
APPLICANT: Smithgall, Molly
                                                                                                                                                                                                                     7.09e+03
38.00
87.50%
75.00%
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9.01e+03
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63.64%
63.64%
3.44%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Primer
                                                                                                                                                ; ORGANISM: Homo sapiens
US-09-801-274-943
                                                                                                                                                                                                                                                        Percent Similarity:
Best Local Similarity:
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Best Local Similarity:
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US-10-125-635A-370/c
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                                                                                                                               TYPE: DNA
                                                                                                                                                                                                                                                                                              Query Match:
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DB:
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                                                                                                                                                                                                                     Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ed. No.:
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APPLICANT: Sleath, Paul R.
APPLICANT: Sleath, Paul R.
APPLICANT: Mossman, Lawrence
APPLICANT: Spies, A. Gregory
APPLICANT: Spies, A. Gregory
APPLICANT: Boydston, Jeremy
ITILE OF INVENTION: COMPOSITIONS AND METHODS FOR WT1
TILLE OF INVENTION: SPECIFIC INMUNOTHERAPY
FILE REFERENCE: 210121.465C5
CURRENT APPLICATION NUMBER: US/09/938,864
CURRENT PILING DATE: 2001-08-24
NUMBER OF SEQ ID NOS: 413
SOFTWARE: FastSEQ for Windows Version 3.0
IENGTH: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: CALSCIP.
APPLICANT: MOMEIL, Patricia D.
APPLICANT: MOMEIL, Patricia D.
APPLICANT: MONEIL, Patricia D.
APPLICANT: Smithgall, Molly
APPLICANT: Solitor, Thomas S.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Sleath, Paul
TITLE OF INVENTION: SPECIFIC IMMUNOTHERAPY
FILE REFERENCE: 210121,465C4
CURRENT APPLICATION NUMBER: US/09/785,019
CURRENT FILICATION NUMBER: 2001-02-15
NUMBER OF SEQ ID NOS: 376
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 370
LENGTH: 38
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0 0 0 0
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Mismatches:
Indels:
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Mismatches:
Indels:
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Matches:
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Matches:
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                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 9.01e+03
38.00
63.64%
63.64%
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63.648
63.648
3.448
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Moulton, Gus
Vedvick, Tho
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Best Local Similarity:
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Best Local Similarity:
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US-09-785-019-370/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match:
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Sequence 11, Application US/09920342
Fatent No. US20020137709A1
GENERAL INFORMATION:
APPLICANT: University of Southern California
APPLICANT: University of Southern California
APPLICANT: Chong, Cheng-Ming
APPLICANT: Randall B.
TITLE OF INVENTION: GENE SILENCING USING MRNA-CDNA HYBRIDS
FILE REFERENCE: 13761-7024
CURRENT FILING DATE: 2002-01-17
PRIOR FILING DATE: 2002-01-17
PRIOR FILING DATE: 2000-08-02
                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR WIT TITLE OF INVENTION: SPECIFIC IMMUNOTHERAPY FILE REFERENCE: 210121.465C6
CURRENT APPLICATION NUMBER: US/10/002,603
UNMBER OF SEQ ID NOS: 413
SOFTWARE: FastSEQ for Windows Version 3.0
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Mismatches:
Indels:
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Matches:
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                       35 GGTCCAGCACCTCCGCCCGCAACGCCCCCCAG 3
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                     ; Sequence 370, Application US/10002603; Publication No. US20030095971A1; GENERAL INFORMATION:
                                                                                                                                                                                APPLICANT: Gaiger, Alexander APPLICANT: McNeill, Patricia D. APPLICANT: Smithgall, Molly APPLICANT: Moulton, Gus APPLICANT: Wodvick, Thomas S. APPLICANT: Yedvick, Paul R. APPLICANT: Mossman, Sally APPLICANT: Evans, Lawrence APPLICANT: Spies, A. Gregory APPLICANT: Boydston, Jeremy APPLICANT: Boydston, Jeremy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; FEATURE:
; OTHER INFORMATION: bc12 primer
US-09-920-342-11
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
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63.64%
63.64%
3.44%
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Best Local Similarity:
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LENGTH: 38
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LENGTH: 24
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TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
FILE REFERENCE: P8017
CURRENT APPLICATION NUMBER: US/10/0667
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PELICATION NUMBER: PCT/US01/00669
PRIOR FILING DATE: 2001-01-30
PRIOR PELICATION NUMBER: PCT/US01/00668
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR PILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-05-23
PRIOR PLICATION NUMBER: US 60/327, 898
PRIOR PILING DATE: 2001-05-30
PRIOR PILING DATE: 2001-05-30
PRIOR PILING DATE: 2001-05-30
PRIOR PILING DATE: 2001-00-09
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Publication No. US20030046717A1
GENERAL INFORMATION:
APPLICANT: Zhang, Jian
TITLE OF INVENTION: HUMAN TESTIS EXPRESSED PATCHED LIKE PROTEIN
FILE REFERRENCE: P80177
CURRENT APPLICATION NUMBER: US/10/060,756A
CURRENT FILING DATE: 2002-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
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Mismatches:
Indels:
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Matches:
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                   Length:
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                     6.64e+03
                                           37.00
100.00%
100.00%
3.35%
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75.00%
75.00%
3.35%
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Best Local Similarity:
Query Match:
                                                   Percent Similarity:
Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                      RESULT 16
US-10-060-756A-2700
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Alignment Scores:
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30 CAGGACAGTGGGCGRTGGGTAGGC 7
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                                         RESULT 19
US-09-874-503-34/c
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Sequence 1740, Application US/09801274

Patent No. US20020031319A1

Patent No. US20020031319A1

GENERAL INFORMATION:

APPLICANT: Cargill, Michele

APPLICANT: Lander, Eric S.

TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS

TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS

TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS

CURRENT FILING DATE: 2009-001

CURRENT FILING DATE: 2001-03-07

PRIOR FILING DATE: 2000-03-07

PRIOR FILING DATE: 2000-03-07

PRIOR FILING DATE: 2000-03-07

PRIOR FILING DATE: 2000-05-22

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 1740

LENGTH: 31
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Matches:
Conservative:
Mismatches:
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Matches:
Conservative:
Mismatches:
Indels:
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                                                      PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR PLING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR PLING DATE: 2001-05-23
PRIOR PLING DATE: 2001-06-23
PRIOR APPLICATION NUMBER: US 60/327,898
PRIOR PILING DATE: 2001-10-09
PRIOR APPLICATION NUMBER: US 60/327,898
PRIOR PILING DATE: 2001-10-09
PRIOR APPLICATION NUMBER: ASO1-10-09
PRIOR APPLICATION NUMBER: US 60/327,898
PRIOR APPLICATION NUMBER: ASO1-10-09
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                      PPLICATION NUMBER: PCT/US01/00664
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37.00
87.50%
75.00%
3.35%
                                           ILING DATE: 2001-01-30
FILING DATE: 2001-01-30
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75.00%
75.00%
3.35%
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ORGANISM: Homo sapiens
18-10-060-756A-2701
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Percent Similarity:
Best Local Similarity:
Query Match:
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Best Local Similarity:
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LENGTH: 25
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190 GlnAspAsnGlyGlyTrpValGly 197

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APPLICANT: WOOLWILLIAM 1.

APPLICANT: WOOLWILLIAM 1.

TITLE OF INVENTION: IL-17 HOMOLOGOUS POLYPEPTIDES AND THERAPEUTIC USES THEREOF
FILE REFERENCE: 19381R1C1P3 (US)
CURRENT APPLICATION NUMBER: US 60/253, 646
PRIOR APPLICATION NUMBER: US 60/244,072
PRIOR APPLICATION NUMBER: US 60/244,072
PRIOR APPLICATION NUMBER: US 60/242,837
PRIOR APPLICATION NUMBER: US 60/242,837
PRIOR FILING DATE: 2000-01-0-26
PRIOR FILING DATE: 2000-01-0-26
PRIOR FILING DATE: 2000-01-11
PRIOR FILING DATE: 2000-01-11
PRIOR FILING DATE: 2000-01-11
PRIOR FILING DATE: 2000-03-21
PRIOR PRILCATION NUMBER: US 60/13,807
PRIOR PELICATION NUMBER: US 60/132,807
PRIOR PELICATION NUMBER: US 60/134,887
PRIOR PELICATION NUMBER: US 60/134,287
PRIOR PELICATION NUMBER: US 60/134,287
PRIOR PELICATION NUMBER: US 60/134,287
PRIOR PELICATION NUMBER: US 60/131,022
PRIOR APPLICATION NUMBER: US 60/131,022
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APPLICATION NUMBER: US PCT/US01/06520
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R APPLICATION NUMBER: US 60/130,232

R FILING DATE: 1999-04-21

R FILING DATE: 1998-12-23

R FILING DATE: 1998-12-23

R RFILING DATE: 1998-15-15

R APPLICATION NUMBER: US 60/085,579

R APPLICATION NUMBER: US 09/854,208

R APPLICATION NUMBER: US 09/854,208

R FILING DATE: 2001-05-10

R FILING DATE: 2001-05-20
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R FILING DATE: 2001-03-22
R PILING DATE: 2000-12-20
R FILING DATE: 2000-12-20
R APPLICATION NUMBER: US 09/644,848
R FILING DATE: US 09/644,848
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APPLICATION NUMBER: US 09/380,138
FILING DATE: 1999-08-25
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Sequence 34, Application US/09874503 Patent No. US20020177188A1
                                                                                                                                                                                               Godowski, Paul J.
Grimaldi, J. Christopher
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VanLookeren, Menno
Vandlen, Richard
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Hillan, Kenneth J.
Hymowitz, Sarah G.
Tumas, Daniel
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                                                                                                              Filvaroff, Ellen
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PRIOR FILING DATE: JOUL 09.5

PRIOR PELICATION NUMBER: 09/931836

PRIOR PELING DATE: 2001-08-16

PRIOR FILING DATE: 1999-03-08

PRIOR FILING DATE: 1999-05-14

PRIOR FILING DATE: 1999-05-14

PRIOR FILING DATE: 1999-05-14

PRIOR FILING DATE: 1999-05-14

PRIOR PELICATION NUMBER: PCT/US09/31274

PRIOR PELICATION NUMBER: PCT/US00/04341

PRIOR PLING DATE: 2000-02-18

PRIOR PLING DATE: 2001-03-01

PRIOR PLING DATE: 2001-03-01

PRIOR PLING DATE: 2000-03-18

PRIOR PLING DATE: 2000-03-02

PRIOR PLING DATE: 2000-03-02

PRIOR PLING DATE: 2000-03-02

PRIOR PLING DATE: 2000-03-21

PRIOR PLING DATE: 2000-06-02

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OR FILING DATE: 2001-08-13

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OR FILING DATE: 1999-03-08

OR FILING DATE: 1999-03-08

OR PILING DATE: 1999-03-08

OR PILING DATE: 1999-13-08

OR APPLICATION NUMBER: PCT/US99/10733

OR PILING DATE: 1999-12-30

OR PILING DATE: 1999-05-14

OR FILING DATE: 1999-05-14

OR PILING DATE: 2000-02-18

OR APPLICATION NUMBER: PCT/US00/04341

OR PILING DATE: 2000-02-18
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PRIOR FILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: PCT/US00/30873
PRIOR FILING DATE: 2000-11-10
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PRIOR APPLICATION NUMBER: 09/874503
PRIOR PETLING DATE: 2001-06-05
PRIOR PETLING DATE: 2001-07-18
PRIOR FILING DATE: 2001-07-18
PRIOR FILING DATE: 2001-07-18
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PRIOR PLING DATE: 1998-12-23
PRIOR APPLICATION NUMBER: 60/13023
PRIOR APPLICATION NUMBER: 60/13102
PRIOR PILING DATE: 1999-04-26
PRIOR FILING DATE: 1999-04-26
PRIOR FILING DATE: 1999-05-14
PRIOR FILING DATE: 1999-05-14
PRIOR PRILING DATE: 1999-06-09
PRIOR PLING DATE: 1999-06-09
PRIOR PLING DATE: 1999-06-09
PRIOR APPLICATION NUMBER: 60/130387
PRIOR APPLICATION NUMBER: 60/130387
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PRIOR APPLICATION NUMBER: 09/747259
PRIOR FILING DATE: 2000-12-20
PRIOR APPLICATION TO BE TO THE T
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PRIOR APPLICATION NUMBER: 09/854208
PRIOR FILING DATE: 2001-05-10
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PRIOR APPLICATION NUMBER: 09/644848
PRIOR FILING DATE: 2000-08-22
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FILING DATE: 2000-06-22
APPLICATION NUMBER: 60/242837
FILING DATE: 2000-10-24
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FILING DATE: 2000-10-26
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PRIOR APPLICATION NUMBER: 09/380138
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PRIOR APPLICATION NUMBER: 09/380142
CURRENT FILING DATE: 2001-10-30
PRIOR APPLICATION NUMBER: 60/085579
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APPLICATION NUMBER: 60/175481
FILING DATE: 2000-01-11
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FILING DATE: 2000-03-21
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APPLICATION NUMBER: 60/113621
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TITLE OF INVENTION: IL-17 HOMOLOGOUS POLYPEPTIDES AND THERAPEUTIC USES THEREOF
FILE REFERENCE: P1381R1C1P4(US)
CURRENT APPLICATION UNDBER: US/10/000,157
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Matches:
Conservative:
Mismatches:
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                                                                 PRIOR APPLICATION NUMBER: US PCT/USO0/32678
PRIOR FILING DATE: 2000-12-0
PRIOR FILING DATE: 2000-11-10
PRIOR PELICATION NUMBER: US PCT/USO0/30873
PRIOR PELING DATE: 2000-11-10
PRIOR PELING DATE: 2000-11-10
PRIOR PELING DATE: 2000-62-4
PRIOR APPLICATION NUMBER: US PCT/USO0/15264
PRIOR FILING DATE: 2000-66-02
PRIOR PELING DATE: 2000-68-10
PRIOR PELING DATE: 2000-63-21
PRIOR PELING DATE: 2000-63-10
PRIOR FILING DATE: 2000-03-10
PRIOR PLING DATE: 2000-03-10
PRIOR PELING DATE: 2000-03-18
PRIOR PELING DATE: 2000-03-18
PRIOR PELING DATE: 2000-03-18
PRIOR PELING DATE: 1999-12-30
PRIOR PELING DATE: 1999-12-30
PRIOR APPLICATION NUMBER: US PCT/US99/31274
PRIOR APPLICATION NUMBER: US PCT/US99/31274
PRIOR APPLICATION NUMBER: US PCT/US99/31274
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PRIOR APPLICATION NUMBER: US PCT/US99/05028
PRIOR FILING DATE: 1999-03-08
NUMBER OF SEQ ID NOS: 39
ESEQ ID NO 34
LENGTH: 33
                                       FILING DATE: 2000-12-20
APPLICATION NUMBER: US PCT/US00/32678
                APPLICATION NUMBER: US PCT/US00/34956
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Grimaldi, J.Christopher
Gurney, Austin
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Williams, P.Mickey
Wood, William
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Hillan, Kenneth J.
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APPLICANT: Filvaroff, Ellen
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Tumas, Daniel
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Goddard, Audrey
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Best Local Similarity:
Query Match:
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Yansura, Daniel
VENTION: IL-17 HOMOLOGOUS POLYPEPTIDES AND THERAPEUTIC USES THEREOF
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Mismatches:
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Matches:
                              PRIOR FILING DATE: 2000-03-02

PRIOR PLILING DATE: 2000-03-02

PRIOR APPLICATION NUMBER: US 60/191,007

PRIOR FILING DATE: 2000-03-21

PRIOR PLILING DATE: 2000-03-21

PRIOR PLILING DATE: 2000-03-21

PRIOR PLILING DATE: 2000-06-02

PRIOR FILING DATE: 2000-06-02

PRIOR FILING DATE: 2000-06-22

PRIOR PLILING DATE: 2000-06-22

PRIOR PLILING DATE: 2000-08-22

PRIOR PLILING DATE: 2000-08-22

PRIOR PRILING DATE: 2000-08-22

PRIOR PRILING DATE: 2000-08-24

PRIOR PRILING DATE: 2000-08-24

PRIOR PRILING DATE: 2000-10-24

PRIOR PLILING DATE: 2000-11-10

PRIOR PLILING DATE: 2000-11-10
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CURRENT APPLICATION NUMBER: US/09/908,827
CURRENT FILING DATE: 2001-07-18
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PRIOR APPLICATION NUMBER: 60/085,579
PRIOR FILING DATE: 1998-05-15
PRIOR APPLICATION NUMBER: 60/113,621
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Grimaldi, Christopher
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VanLookeren, Menno
Vandlen, Richard
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Williams, P.Mickey
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APPLICANT: Filvaroff, Ellen
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Hillan, Kenneth
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Best Local Similarity:
Query Match:
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CURRENT APPLICATION NUMBER: US/09/747,259
PRIOR APPLICATION NUMBER: US 09/311,832
PRIOR APPLICATION NUMBER: US 60/172,096
PRIOR FILING DATE: 1999-05-14
PRIOR PAPLICATION NUMBER: US 60/172,096
PRIOR PRIOR DATE: 1999-12-33
PRIOR PELIOR DATE: 1999-12-30
PRIOR FILING DATE: 1999-12-30
PRIOR FILING DATE: 1090-01-11
PRIOR PELICATION NUMBER: US 60/175,481
PRIOR PELICATION NUMBER: US 60/175,481
PRIOR PELICATION NUMBER: PCT/USO0/04341
PRIOR FILING DATE: 2000-02-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Synthetic Oligonucleotide Probe [S-10-000-157-34]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches:
Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-375-514-22 (1-205) x US-10-000-157-34 (1-33)
               PRIOR FILING DATE: 2000-12-01
PRIOR APPLICATION WOMBER: PCI/USOU/318/8
PRIOR APPLICATION UNMBER: PCI/USOU/34956
PRIOR FILING DATE: 2000-12-20
PRIOR FILING DATE: 2001-02-28
PRIOR FILING DATE: 2001-06-01
PRIOR FILING DATE: 2001-06-01
PRIOR FILING DATE: 2001-06-01
PRIOR FILING DATE: 2001-06-01
PRIOR APPLICATION NUMBER: PCI/USO1/19692
PRIOR FILING DATE: 2001-06-29
PRIOR FILING DATE: 2001-07-09
NUMBER OF SEQ ID NOS: 39
LENGTH: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels:
PRIOR APPLICATION NUMBER: PCT/US00/32678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 34, Application US/09747259 Publication No. US20030008815A1 GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            56 ThrProHisProAlaAlaSerArg 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25 ACACCCCATCCTGTAGCCACTCGC 2
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Grimaldi, Christopher
                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
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37.00
87.50%
75.00%
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Vandlen, Richard
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Williams, P.Mickey
Wood, William
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Filvaroff, Ellen
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Hillan, Kenneth
Tumas, Daniel
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Goddard, Audrey
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Best Local Similarity:
Query Match:
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US-09-747-259-34/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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APPLICANT:
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APPLICANT:
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Pred. No.:

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OTHER INFORMATION: Synthetic Oligonucleotide Probe
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PRIOR FILING DATE: 1999-12-30
PRIOR PLILING DATE: 2000-02-18
PRIOR FILING DATE: 2000-02-18
PRIOR FILING DATE: 2000-03-01
PRIOR PELING DATE: 2000-03-02
PRIOR PELING DATE: 2000-03-02
PRIOR PILING DATE: 2000-03-02
PRIOR PILING DATE: 2000-03-12
PRIOR PLILING DATE: 2000-03-21
PRIOR PILING DATE: 2000-03-21
PRIOR PLILING DATE: 2000-03-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OR FILING DATE: 2001-02-20
OR APPLICATION NUMBER: 09/816,744
OR FILING DATE: 2001-03-22
OR FILING DATE: 2001-03-22
OR FILING DATE: 2001-05-10
OR FILING DATE: 2001-05-10
OR FILING DATE: 2001-05-10
OR FILING DATE: 2001-05-10
OR PILING DATE: 1999-03-08
OR PRING DATE: 1999-03-08
OR PRING DATE: 1999-05-14
OR FILING DATE: 1999-05-14
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PRIOR APPLICATION NUMBER: PCT/US01/06520
PRIOR FILING DATE: 2001-02-28
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PRIOR APPLICATION NUMBER: PCT/US00/32678
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APPLICATION NUMBER: PCT/USO0/30873
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FILING DATE: 1999-12-30
                                                                                                                                                                                               OR FILING DATE: 1999-06-09
OR APPLICATION NUMBER: 60/172,096
OR FILING DATE: 1999-12-23
OR APPLICATION NUMBER: 60/175,481
OR APPLICATION NUMBER: 60/175,481
OR APPLICATION NUMBER: 60/191,007
OR FILING DATE: 2000-01-11
OR APPLICATION NUMBER: 60/213,807
OR FILING DATE: 2000-06-22
OR APPLICATION NUMBER: 60/242,837
OR FILING DATE: 2000-10-24
OR FILING DATE: 2000-10-24
OR PILING DATE: 2000-10-24
OR FILING DATE: 2000-10-24
OR PILING DATE: 2000-10-26
OR FILING DATE: 1999-05-14
                R APPLICATION NUMBER: 60/130, 232
RR FILING DATE: 1999-04-21
RR APPLICATION NUMBER: 60/131, 022
RR APPLICATION NUMBER: 60/134, 287
RR APPLICATION NUMBER: 60/138, 387
RR FILING DATE: 1999-05-14
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APPLICATION NUMBER: 09/380,142
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APPLICATION NUMBER: 09/644,848
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR FILING DATE: 2000-12-01
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FILING DATE: 1998-12-23
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SEQ ID NO 34
LENGTH: 33
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Sequence 71, Application US/09245802
Sequence 71, Application US/09245802
Patent No. US20010049125A1
GENERAL INFORMATION:
APPLICANT: STEMMER PH.D., WILLEM P.C.
APPLICANT: LIPSHUTZ, ROBERT J.
TITLE OF INVENTION: END-COMPLEMENTARY POLYMERASE REACTION
NUMBER OF SEQUENCES: 136
CORRESPONDENCE ADDRESS: 00000 001100 OVERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: Conservative: Mismatches: Conservative: US-09-375-514-22 (1-205) x US-09-245-802-71 (1-40) ADDRESSEE: TOWNSEND AND TOWNSEND AND CREW STREET: TWO EMBARCADERO CENTER, 8TH FLOOR CITY: SAN FRANCISCO STATE: CALLFORNIA Mismatches: US-09-375-514-22 (1-205) x US-09-908-827-34 (1-33) RAME: MORPHY, ESQ., MATTHEW B. REGISTRATION NUMBER: 39,787
REGISTRATION NUMBER: 1652BJ-015410US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 326-2400
TELEPHONE: (415) 326-2422
INFORMATION FOR SEQ ID NO: 71: Matches: Length: Matches: Length: Indels: Indels: Gaps: 51 SerGlnProGlyHisThrProHisProAla 60 CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/05480
FILING DATE: 18-APR-1996
CLASSIFICATION:
ATTORNEY/AGBNT INFORMATION:
ATTORNEY/AGBNT INFORMATION: APPLICATION NUMBER: US/09/245,802 PRIOR APPLICATION DATA: APPLICATION NUMBER: 08/675,502 COUNTRY: U.S.A.
ZIP: 94111-3834
COMPUTER READABLE FORM:
WEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DO 56 ThrProHisProAlaAlaSerArg 63 25 ACACCCCATCCTGTAGCCACTCGC 2 1.21e+04 37.00 70.00% 70.00% 3.35% SEQUENCE CHARACTERISTICS: 37.00 87.50% 75.00% 3.35% LENGTH: 40 base pairs TYPE: nucleic acid STRANDEDNESS: double circular Best Local Similarity: Percent Similarity: Best Local Similarity: Query Match: CLASSIFICATION: Percent Similarity: FILING DATE: FILING DATE Alignment Scores: US-09-245-802-71/c TOPOLOGY: US-09-245-802-71 Query Match: RESULT 23 Score: δλ q

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APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MAT.
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
TITLE OF INVENTION: Myosin-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
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Mismatches:
Indels:
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Mismatches:
Indels:
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                                                          ; OTHER INFORMATION: Antisense Oligonucleotide US-09-734-847A-45
                                                                                                                                                   Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Farr, Spencer B.
APPLICANT: Pickett, Gavin G.
APPLICANT: Dickett, Gavin G.
APPLICANT: Dunn, II, Robert Thomas
TITLE OF INVENTION: CANINE TOXICITY GENES
TILLE NEFERENCE: 40074200200
CURRENT. APPLICATION NUMBER: US/09/911, 904
CURRENT FILING DATE: 2002-04-09
PRIOR PILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 386
SOFTWARE: FEASTSEQ for Windows Version 4.0
SEQ ID NO 37
LENGTH: 25
                                                                                                                                                                                                                                                          Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 4774, Application US/09866108
Patent No. US20020048800A1
                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-911-904-37/c
; Sequence 37, Application US/09911904
; Publication No. US20030096234A1
; GENERAL INFORMATION:
                    ORGANISM: Artificial Sequence
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36.00
100.008
83.33
3.268
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85.71%
71.43%
3.26%
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Canis familiaris
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Best Local Similarity:
Query Match:
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Best Local Similarity:
Query Match:
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                                                                                                                      Alignment Scores:
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                                                                                                                                        Pred. No.:
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APPLICANT: Ereir, Susan
APPLICANT: McKay, Robert
APPLICANT: Maria, James G.
TTLLE OF INVENTION: Alteration of Cellular Behavior by Antisense Modulation of mRNA
FILE REFERENCE: ISPH-0524
CURRENT FILING DATE: 1200-12-12
PRIOR FILING DATE: 1998-10-07
PRIOR APPLICATION NUMBER: 09/167,921
PRIOR FILING DATE: 1999-03-26
                                                                                                                                   APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, Bentet P.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Alang, Oingoing
TITLE OF INVENTION: Antisense Modulation of bcl-x Expression
FILE REFERENCE: ISPH-0528
CURRENT APPLICATION NUMBER: US/09/734,846
CURRENT FILING DATE: 2000-12-12
PRIOR APPLICATION NUMBER: 09/277,020
PRIOR APPLICATION NUMBER: 09/277,020
PRIOR APPLICATION NUMBER: 09/167,921
PRIOR APPLICATION NUMBER: 09/167,921
PRIOR FILING DATE: 1998-10-07
PRIOR FILING DATE: 1999-06-02
NUMBER OF SEQ ID NOS: 74
SCENTHARE: PATENTIN VOIL STAND NOS: 74
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Antisense Oligonucleotide 3S-09-734-846-34
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Matches:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 45
LENGTH: 20
TYPE: DNA
37 AGCCAGCCCGACACCCGCCAACACCCGCT 8
                                                          US-09-734-846-34/c
; Sequence 34, Application US/09734846; Patent No. US20010007025A1
; BRIERAL INFORMATION:
; APPLICANT: Bennett, C. Frank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 25
US-09-734-847A-45/c
; Sequence 45, Application US/09734847A
; Patent No. US20020049173A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Bennett, C. Frank
APPLICANT: Crooke, Stanley T.
APPLICANT: Manoharan, Muthiah
APPLICANT: Myatt, Jacqueline R.
APPLICANT: Baker, Brenda F.
APPLICANT: Monia, Brett P.
APPLICANT: Freir, Susan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6.78e + 03
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100.008
83.338
3.268
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Best Local Similarity:
Query Match:
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                                        RESULT 24
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APPLICANT: PENN, Sharron G.
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
TITLE OF INVENTION: WOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REPERENCE: ABOMICA-7
CURRENT PILING DATE: 2001-05-25
PRIOR PILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-06-40
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: NCS 60/236, 359
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR PRILICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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Matches:
Conservative:
Mismatches:
Indels:
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PRIOR APPLICATION NUMBER: PCT/USO1/00666
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00664
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-02-05
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APPLICANT: JI, Yonggang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Homo sapiens US-09-866-108-4775
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Query Match:
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APPLICANT: Ji, Yonggang
APPLICANT: Ji, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MAT.
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
PILE REFERENCE: AEOMICA-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             00 10 0 22
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Mismatches:
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Matches:
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR PRILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
                                               PRIOR FILING DATE: 2000-10-10
PRIOR PELLING DATE: 2000-10-04
PRIOR PELLING DATE: 2000-10-04
PRIOR PELLING DATE: 2000-10-07
PRIOR PELLING DATE: 2000-09-27
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PELLING DATE: 2001-01-30
PRI
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                                    FILING DATE: 2000-05-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Percent Similarity:
Best Local Similarity:
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APPLICATION NUMBER: PCT/US01/00668
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US-09-866-108-4778/c
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APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
TITLE OF INVENTION: Mark
TITLE OF INVENTION: USCO9/866,108
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,666
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PRILIKG DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR APPLICATION NUMBER: PCT/US01/00667
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Conservative:
Mismatches:
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                                          FRIOR FILING DATE: 2001-01-30
PRIOR PELICATION NUMBER: PCT/US01/00663
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomica Sequence Listing Engine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length:
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                           APPLICATION NUMBER: PCT/US01/00669 FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00665
FILING DATE: 2001-01-30
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Patent No. US20020048800al
GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G,
APPLICANT: HANZEL, David K.
FILING DATE: 2001-01-30
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83.33%
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3.26%
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CORGANISM: Homo sapiens
US-09-866-108-4776
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LENGTH: 25
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APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
TITLE OF INVERTION: MYSON-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-26
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-06-26
PRIOR PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR DATE: 2001-01-30
PRIOR PRIING DATE: 2001-01-30
PRIOR PRIOR DATE: 2001-01-30
PRIOR PRIOR PRIING DATE: 2001-01-30
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PRIOR PRIOR DATE: 2001-01-30
PRIOR PRIOR DATE: 2001-01-30
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Mismatches:
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Matches:
PRIOR FILING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PELICATION NUMBER: PCT/USO1/00662
PRIOR PELING DATE: 2001-01-30
PRIOR PELING DATE: 2001-01-30
PRIOR PELING DATE: 2001-01-30
PRIOR PELING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00670
PRIOR APPLICATION NUMBER: CS/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR PELING DATE: 2000-09-21
PRIOR PELING DATE: 2000-09-21
PRIOR PELING DATE: 2000-09-25
PRIOR PELING DATE: 2001-02-05
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                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 4777
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps:
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Patent No. US20020048800A1
GENERAL INFORMATION:
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HANZEL, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22 GGACATGCTCCTCATCCA 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-4777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: GU, Yizhong
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Best Local Similarity:
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Percent Similarity:
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STREAM. INFORMATION:

APPLICANT: GU, Yizhong

APPLICANT: JENN, Sharron G.

APPLICANT: JENN, Sharron G.

APPLICANT: HENN, Sharron G.

APPLICANT: HENN, Sharron G.

APPLICANT: HENN, Sharron G.

APPLICANT: HENN, Wensheng

APPLICANT: SHANNON, Mark

TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE

TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE

TITLE OF INVENTION: WOMBER: US 60/207,456

PRIOR PAPLICATION NUMBER: US 60/207,456

PRIOR PAPLICATION NUMBER: GB 24.263.6

PRIOR PELING DATE: 2001-01-04

PRIOR PELING DATE: 2001-01-04

PRIOR PELING DATE: 2001-01-04

PRIOR PELING DATE: 2001-01-04

PRIOR PELING DATE: 2001-01-130

PRIOR PELING DATE: 2001-01-30

PRIOR PELING DATE: 2001-01-30
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Matches:
Conservative:
Mismatches:
Indels:
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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-21
PRIOR PILING DATE: 2000-09-21
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Accomica Sequence Listing Engine
SEQ ID NO 4778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 4779, Application US/09866108 Patent No. US20020048800A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8.82e+03
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83.33%
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3.26%
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US-09-866-108-4778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Percent Similarity:
Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-866-108-4779/C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 32
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PRIOR PRILATION NUMBER: 050/234,687

PRIOR PAPLICATION NUMBER: 050/00-09-18

PROFINAR: AMOUNT SEQUENCE LISTING Engine

SOCTION OF 050-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19-05-19
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APPLICANT: New York Blood Center
TITLE OF INVENTION NUMBER: US/09/49/145
CURRENT APPLICATION NUMBER: US/09/949/145
CURRENT FILING DATE: 2000-09-07
PRIOR FILING DATE: 2000-09-07
PRIOR FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 77
SEQ ID NO 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Description of Artificial Sequence:
; OTHER INFORMATION: Oligonucleotides, linkages between positions
; OTHER INFORMATION: to 20 are phosphorothioate, linkages between
; OTHER INFORMATION: positions 20 to 28 are N3' to N5'
; OTHER INFORMATION: phosphoramidates
; OTHER INFORMATION: position 28 is modified by a 3'-aminodeoxyribosyl
; OTHER INFORMATION: residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 11, Application US/09817387
Patent No. US20010039263A1
GENERAL INFORMATION:
APPLICANT: Max-Delbruck-Centrum fur Molekulare Medizin
APPLICANT: Max-Delbruck-Centrum fur Molekulare Medizin
TITLE OF INVENTION: Chimeric Oligonucleotides and the Use Thereof
FILE REFERENCE: 101195-24
CURRENT FILING DATE: 2001-03-26
FRIOR APPLICATION NUMBER: US/09/817,387
PRIOR FILING DATE: 1997-05-02
                                                                                                           001022
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Matches:
Conservative:
Mismatches:
Indels:
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Mismatches:
Indels:
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                                                                        8.82e+03
36.00
83.33&
83.33&
3.26&
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ORGANISM: Artificial Sequence
FEATURE:
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SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 11
LENGTH: 28
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                                                                                                                                                                 Percent Similarity:
Best Local Similarity:
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Best Local Similarity:
Query Match:
               US-09-866-108-4781
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                                                                           Alignment Scores:
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                                                                                                     Pred. No.:
                                                                                                                                           Score:
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REPERENCE: ABOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PLICATION NUMBER: GB 24263.6
PRIOR PLING DATE: 2000-05-26
PRIOR PLING DATE: 2000-010-04
PRIOR PLING DATE: 2000-010-04
PRIOR PLING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
                                                                                                                                                                                                                                                                                                         Conservative:
Mismatches:
Indels:
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Matches:
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OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00667
OR APPLICATION NUMBER: PCT/US01/00664
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00669
OR APPLICATION NUMBER: PCT/US01/00669
OR PILING DATE: 2001-01-30
OR PILING DATE: 2001-01-30
OR PILING DATE: 2001-01-30
OR PILING DATE: 2001-01-30
OR FILING DATE: 2001-01-30
SOFTWARE: Aeomica Sequence Listing Engine SEQ ID NO 4780
LENGTH: 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: PCT/US01/00666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: PCT/USO1/00662
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/USO1/00661
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/USO1/00670
FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 4781, Application US/09866108
Patent No. US200200488800A1
GENERAL INFORMATION:
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36.00
83.33%
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PENN, Sharron G.
HANZEL, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19 GGACATGCTCCTCATCCA 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RANK, David R. CHEN, Wensheng
                                                                                                                        ; ORGANISM: Homo sapiens US-09-866-108-4780
                                                                                                                                                                                                                                                                                                         Percent Similarity:
Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                Alignment Scores:
Pred. No.:
                                                                                              TYPE: DNA
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Sequence 8, Application US/10245206
Publication No. US20030073122A1
GENERAL INFORMATION:
APPLICANT: Nerenberg, Michael I.
APPLICANT: Canter, David M.
APPLICANT: Radtkey, Ray R.
TITLE OF INVENTION: METHODS FOR DETERMINATION OF SINGLE
TITLE OF INVENTION: BIOELECTRIC MICROCHIP
                                                                                                                                                    00100
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Matches:
Conservative:
Mismatches:
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Mismatches:
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                                                                                                                                                  Length:
Matches:
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CURRENT APPLICATION NUMBER: US/10/245,206

CURRENT FILING DATE: 2002-09-16

PRIOR APPLICATION NUMBER: US/09/291,129

PRIOR FILING DATE: 1999-04-12

PRIOR FILING DATE: 1999-04-12

PRIOR FILING DATE: 1998-02-25

PRIOR FILING DATE: 1998-02-25

PRIOR FILING DATE: 1997-12-05

PRIOR FILING DATE: 1997-12-05

PRIOR FILING DATE: 1997-12-05

PRIOR FILING DATE: 1997-12-05

PRIOR FILING DATE: 1997-05-20

PRIOR FILING DATE: 1996-09-06

PRIOR FILING DATE: 1996-09-06

PRIOR FILING DATE: 1996-09-06

PRIOR FILING DATE: 1996-09-06

PRIOR FILING DATE: 1996-09-07

PRIOR PRIOR DATE: 1996-09-07

PRIOR PRIOR DATE: 1994-09-09

PRIOR FILING DATE: 1994-09-09

PRIOR PRIOR FILING DATE: 1994-09-09

PRIOR PRIOR PRIOR DATE: 1994-09-09
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SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                       1.05e+04
36.00
83.33%
83.33%
3.26%
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36.00
75.00%
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                                                ; ORGANISM: Mus musculus US-09-949-145-28
                                                                                                                                                                                             Percent Similarity:
Best Local Similarity:
Query Match:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-245-206-8/c
                                                                                                                   Alignment Scores:
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LENGTH: 29
TYPE: DNA
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ProGlyAlaAlaProAlaProGly 47

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Sequence 1215, Application US/09801274

Sequence 1215, Application US/09801274

Batent No. US20020032319A1

GENERAL INFORMATION:

APPLICANT: Cargill, Michele

APPLICANT: Lander, Eric S.

TILLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS

FILE REFERENCE: 2825.2009-001

CURRENT APPLICATION NUMBER: US/09/801,274

CURRENT FILING DATE: 2001-03-07

PRIOR PILING DATE: 2000-03-07

PRIOR PILING DATE: 2000-05-22

NUMBER OF SEQ ID NOS: 1802
                                                                                     Sequence 8, Application US/09291129

Patent No. US20010014449A1

GENERAL INFORMATION:

APPLICANT: Merenberg, Michael I.

APPLICANT: Canter, David M.

APPLICANT: Canter, David M.

APPLICANT: Canter, David M.

APPLICANT: Canter, David M.

TITLE OF INVENTION: BIOELECTRIC MICROCHIP

TITLE OF INVENTION: BIOELECTRIC MICROCHIP

FILE REFERENCE: 240/240

CURRENT APPLICATION NUMBER: US 09/030,156

EARLIER FILING DATE: 1999-04-12

EARLIER FILING DATE: 1997-12-05

EARLIER FILING DATE: 1997-12-05

EARLIER FILING DATE: 1997-12-05

EARLIER FILING DATE: 1996-10-04

EARLIER FILING DATE: 1996-10-04

EARLIER FILING DATE: 1996-09-07

EARLIER FILING DATE: 1996-09-07

EARLIER FILING DATE: 1994-09-07

EARLIER FILING DATE: 1994-09-07

EARLIER FILING DATE: 1994-07-07

EARLIER FILING DATE: 1993-11-01

WIMPER OF SED IN NOWER: US 08/146,504
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Mismatches:
Indels:
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Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 8 LENGTH: 30
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30 CCAGGTGGAGCACCCAGGCCTGGA 7
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36.00
75.00%
75.00%
3.26%
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Best Local Similarity:
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ORGANISM: human
                                                                        US-09-291-129-8/c
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Sequence 23, Application US/10000512
Patent No. US20020164699A1
GENERAL INFORMATION:
APPLICANT: Shimkets, Richard A
APPLICANT: Fernandes, Elma
TITLE OF INFURION: POLYPEPTIDES AND POLYNUCLEOTIDES ENCODING SAME
FILE REFERENCE: 15966-55E
CURRENT APPLICATION NUMBER: US/10/000,512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence: PCR primer; OTHER INFORMATION: PDM-251
US-09-854-356-15
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APPLICANT: Cheeven, Dirk
APPLICANT: Cheeven, Dirk
APPLICANT: Cheeven, Dirk
APPLICANT: SmithKline Beecham Biologicals S. A.
TITLE OF INVENTION: HER-2/neu Fusion Proteins
FILE REFERENCE: 014058-0099H30pc
CURRENT APPLICATION NUMBER: US/09/854,356
CURRENT FILING DATE: 2001-05-09
PRIOR APPLICATION NUMBER: US 09/493,480
PRIOR PILING DATE: 2000-01-28
PRIOR FILING DATE: 1999-01-29
                                                                                                                                     Length:
Matches:
Conservative:
Mismatches:
Indels:
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Mismatches:
Indels:
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Matches:
SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 1215
LENGTH: 31
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Patent No. US20020177567A1
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                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
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36.00
100.00%
50.00%
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36.00
100.008
100.008
3.268
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SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 15
LENGTH: 32
                                                     TYPE: DNA ORGANISM: Homo sapiens
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Percent Similarity:
Best Local Similarity:
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Best Local Similarity:
Query Match:
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S-09-854-356-15/c
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Sequence 31, Application US/09473872
Fatent No. US20020064876A1
GENERAL INFORMATION:
APPLICAMT: Yoon, Kyongqeun
TITLE OF INVENTION: No. US20020064876A1e1 Gene Therapy Methods For The Treatment
TITLE OF INVENTION: Disorders
FILE REPERENCE: JEFF-YOOJ
CURRENT APPLICATION NUMBER: US/09/473,872
CURRENT APPLICATION NUMBER: 1999-12-28
NUMBER OF SEQ ID NOS: 68
SSOFTWARE: Patentin Ver. 2.1
SSOFTWARE: Patentin Ver. 2.1
LENGTH: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COL7Al gene
                                                                                                                                                                                                                             ; OTHER INFORMATION: Description of Artificial Sequence: PCR PRIMER US-10-000-512-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: fragment from a normal human
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Matches:
Conservative:
Mismatches:
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Mismatches:
Indels:
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Matches:
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CURRENT FILING DATE: 2001-10-23
PRIOR APPLICATION NUMBER: 09/619,252
PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: 60/167,785
PRIOR FILING DATE: 1999-11-29
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.1
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Patent No. US20020124285A1

GENERAL INFORMATION:

APPLICANT: Conner, Timothy

APPLICANT: Dubois, Patrice

APPLICANT: Malven; Marianne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                1.31e+04
36.00
100.008
85.718
3.268
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36.00
60.00%
60.00%
3.26%
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Best Local Similarity:
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US-09-894-633A-78
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US-09-473-872-31
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                                                                                                                                  SEQ ID NO 23
LENGTH: 35
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DB:
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APPLICANT: RANK, David K.

APPLICANT: RANK, David R.

APPLICANT: CHEN, Wensheng

TITLE OF INVENTION: MAINT

TITLE OF INVENTION: MAIOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE

FILE REFERENCE: ADOMICA-7

CURRENT APPLICATION NUMBER: US/09/866,108

CURRENT FILING DATE: 2001-05-25

PRIOR FILING DATE: 2000-05-26

PRIOR FILING DATE: 2000-10-04

PRIOR FILING DATE: 2000-10-04

PRIOR FILING DATE: 2000-10-04

PRIOR FILING DATE: 2000-10-04

PRIOR FILING DATE: 2000-10-04
                                               Sequence 67, Application US/09735363A
Patent No. US20010041681A1
GENERAL INFORMATION
APPLICANT: Fillon, Mario
APPLICANT: Fillon, Nigel
TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides
FILE REPERENCE: 02811-0181
CURRENT FILLOS DATE: 1090-12-12
CURRENT FILLOS DATE: 1999-12-13
PRIOR FILLING DATE: 1999-12-13
PRIOR FILLING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 87
SOFTWARE: PATENTI VERSION 3.0
SOFTWARE: PATENTI VERSION 3.0
SOFTWARE: PATENTI VERSION 3.0
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Mismatches:
Indels:
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Matches:
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PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
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Patent No. US20020048800A1
                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Artificial Sèquence
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35.50
70.00%
60.00%
3.22%
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APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity:
Query Match:
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                                         US-09-735-363A-67
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PRIOR
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APPLICANT: Masucci, James
TITLE OF INVENTION: PLANT REGULATORY SEQUENCES FOR SELECTIVE CONTROL OF GENE EXPRESSI
FILE REPERENCE: 38-21(15856)84,633A
CURRENT PEPLICATION NUMBER: US/09/894,633A
CURRENT FILING DATE: 2001-06-28
PRIOR FILING DATE: 2000-06-28
PRIOR FILING DATE: 2000-06-28
PRIOR FILING DATE: 2000-06-28
NUMBER OF SEQ ID NOS: 111
SOFFWARE: Patentin version 3.0
SEQ ID NO 78
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APPLICANT: Filion, Mario
APPLICANT: Philip, Nigel
TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides
FILE REFERENCE: 02811-0181
CURRENT FILING DATE: 2000-12-12
PRIOR FILING DATE: 1999-12-13
PRIOR FILING DATE: 1999-12-13
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 87
SOFTWARE: PATENTIN VERSION 3.0
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Matches:
Conservative:
Mismatches:
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Matches:
Conservative:
Mismatches:
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OTHER INFORMATION: synthetic primer sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Synthetic Oligonucleotide US-09-735-363A-2
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                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: artificial sequence
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35.50
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3.22%
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36.00
77.78%
77.78%
3.26%
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Best Local Similarity:
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Query Match:
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LENGTH: 27
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APPLICANT: SHANNO, MARK
TITLE OF INVERTION, MACK
TITLE OF INVERTION, MACK
TITLE OF INVERTION MACSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PELLING DATE: 2000-10-04
PRIOR FILING DATE: 2000-09-27
PRIOR PELLING DATE: 2000-09-27
PRIOR PELLING DATE: 2001-01-30
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Matches:
Conservative:
Mismatches:
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                                                   PRIOR APPLICATION NUMBER: 15.7.
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomica Sequence Listing Engine
SEQ ID NO 4466
LENGTH: 25
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                                  FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
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APPLICATION NUMBER: PCT/US01/00668
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
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APPLICATION NUMBER: PCT/US01/00662
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APPLICATION NUMBER: PCT/US01/00661
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Patent No. US20020048800A1
GENERAL INFORMATION:
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CORGANISM: Homo sapiens
US-09-866-108-4466
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APPLICANT: CHEN, DAVIG R.
APPLICANT: CHEN, Wensheng
TILLE APPLICANT: CHEN, Wensheng
TILLE OF INVERTION: MYSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: ACONICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PRILING DATE: 2000-06-26
PRIOR FILING DATE: 2000-06-26
PRIOR FILING DATE: 2000-06-26
PRIOR FILING DATE: 2001-01-30
PRIOR PRILING DATE: 2001-01-30
PRIOR PRILING DATE: 2001-01-30
PRIOR PRILING DATE: 2001-01-30
PRIOR PRILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00665
PRIOR APPLICATION NUMBER: PCT/US01/00666
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Mismatches:
Indels:
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                                           PRIOR FILING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR PILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PRILING DATE: 2001-05-65
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Abomica Sequence Listing Engine SEQ ID NO 4465
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
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FILING DATE: 2001-01-30
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Patent No. US20020048800A1
GENERAL INFORMATION:
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CORGANISM: Homo sapiens
US-09-866-108-4465
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Best Local Similarity:
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; ORGANISM: Homo sapiens US-09-866-108-4469
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APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MAY.
TITLE OF INVENTION: WYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-05-26
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Matches:
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PRIOR PELING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PILING DATE: 2000-10-04
PRIOR PILING DATE: 2000-09-27
PRIOR PELING DATE: 2000-09-27
PRIOR PLICATION NUMBER: PCT/USO1/00667
PRIOR PELING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR PELING DATE: 2001-01-30
PRIOR PELING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PELING DATE: 2000-09-21
PRIOR PELING DATE: 2000-09-21
PRIOR PELING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomica Sequence Listing Engine
SEQ ID NO 4467
LENGTH: 25
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Patent No. US20020048800A1
GENERAL INFORMATION:
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US-09-866-108-4467
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Query Match:
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APPLICANT: GU, YIZhong
APPLICANT: GU, YIZhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INFURTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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Matches:
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CURRENT PELLING NUMBER: US/09/866,108
CURRENT PELLING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-06-6
PRIOR PILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-07
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/US01/00665
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PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-31, FRIOR PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
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LENGTH: 25
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PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR PLING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
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Sequence 4469, Application US/09866108;
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; SEQ ID NO 4468
; LENGTH: 25
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-4468
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Maximum DB seq length: 40
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Match
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Database

Sequence Sequence

PCT-US95-10973A-80 PCT-US95-10973A-81 US-07-990-965-2 US-08-340-045-7

Sequence 1

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Result

Sequence

Sequence 1, Appli Sequence 1, Appli Sequence 6, Appl Sequence 6, Appl Sequence 48, Appl Sequence 48, Appl Sequence 48, Appl Sequence 48, Appl Sequence 33, Appl Sequence 48, Appl Sequence 48, Appl Sequence 59, Appl Sequence 60, Appl Sequence 60, Appl Sequence 75, Appl Sequence 75, Appl Sequence 75, Appl Sequence 75, Appl Sequence 75, Appl Sequence 76, Appl Sequence 77, Appl Sequence 31, Appl	Sequence 32, Appl
1 US-08-038-071-1 1 US-08-038-071-1 5 PCT-US92-0265-1 6 55498600-2 1 US-08-019-14 1 US-08-091-219-14 1 US-08-091-219-14 1 US-08-091-219-14 1 US-08-091-219-14 1 US-08-151-168-48 1 US-08-151-178-48 1 US-08-179-74-68 1 US-08-179-74-72 1 US-08-179-130-73 1 US-08-179-130-73 1 US-08-179-130-75 1 US-08-179-170-75 1 US-08-1	PCT-US96-11786-32
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3 195 3 105-08-871-302A-7 Sequence 27, 318 11 US-07-37-33-22 Sequence 27, 318 1 US-07-372-35 Sequence 27, 318 1 US-07-372-35 Sequence 27, 318 1 US-07-372-35 Sequence 27, 318 1 US-07-373-131-29 Sequence 27, 318 1 US-08-345-704-19 Sequence 19, 318 1 US-08-345-704-14 Sequence 19, 318 1 US-08-345-705-14 Sequence 19, 318	5.1 5.2 4 05-09-338-705-37 Sequence 37,

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1 US-08-199-507B-49 1 US-08-199-507B-49 1 US-08-421-684-4 2 US-08-421-684-4 4 US-09-440-675-502-4 4 US-09-440-675-502-4 4 US-09-675-502-1 3 US-08-240-730-1 3 US-08-240-730-1 2 US-08-240-730-1 2 US-08-240-730-1 2 US-08-240-730-1 3 US-08-240-730-218 3 US-09-080-285-77 2 US-08-689-255-24 4 US-09-109-663-72 4 US-09-109-689-255-24 4 US-09-109-109-683-72 4 US-09-109-109-69-72 6 US-09-109-109-69-72 6 US-09-109-109-69-72 6 US-09-109-109-69-73 6 US-09-109-109-69-73 6 US-09-109-109-69-73 6 US-09-109-109-69-73 6 US-09-109-109-73 6 US-09-109-73 6 US-09-109-
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444																																				2	/sn 11	Ē	rnompson G. Draper				:88:	& Lyon		_	
9.50																																				-15	Sequence 15, Application US/U/93042. Patent No. 5750390	DRMATION:	APPLICANT: James D. APPLICANT: Kenneth G	INVENTION:	INVENTION:	F INVENTION:	NCE ADDI	EE: Lyon & 611 West	Los Angeles	California : USA	Ξ
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APPLICANT: James D. Thompson
APPLICANT: Fenneth G. Draper
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: TREATMENT OF DISEASES CAUSED
TITLE OF INVENTION: BY EXPRESSION OF THE BCL-2
TITLE OF INVENTION: GENE
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM COMPATIBLE
OPPRATING SYZEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/936,421
FILING DATE: 19920826
CLASSIFICATION: 435
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
APPLICATION NUMBER:
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SYSTEM: IBM P.C. DOS (Version 5.0)
WordPerfect (Version 5.1)
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Matches:
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APPLICATION NUMBER: US/07/936,421
FILING DATE: 19920826
FILING DATE: 19970826
FILING PAPPLICATION: 435
PRIOR APPLICATION DATA:
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611 West Sixth Street
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                                                                                                                                                                                                                                                                    ATTORNEY AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 197/;
TELEPCOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEX: (7-3510
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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CITY: Los Angeles
GTATE: California
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OPERATING SYSTEM:
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US-07-936-421-15
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Best Local Similarity:
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                                                                                                                                                                                                                                                       FILING DATE:
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ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C. STREET: 1100 New York Avenue, N.W., Suite 600
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; Patent No. 5789201
; CENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Genes Coding For Bcl-y, a Bcl-2
TITLE OF INVENTION: Homologue
NUMBER OF SEQUENCES: 53
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative:
Mismatches:
Indels:
PRIOR APPLICATION DATA: including application PRIOR APPLICATION DATA: described below: APPLICATION NUMBER: FILING DATE:
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NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32327
REFERENCE/DOCKET NUMBER: 197/243
TELECOMMUNICATION INFORMATION:
TELEPRX: (213) 955-0440
TELEPRX: 67-3510
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
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CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
RABE: Esmond, Robert W.
REGISTRATION NUMBER: 32,893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 1483
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2540
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                     164
57.00
100.00%
100.00%
5.16%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                          TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Washington
                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: linear
US-07-936-421-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: both
                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                           Percent Similarity:
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MOLECULE TYPE:

24 TGGATHCARGAYAAYGGNGGNTGG 1

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Sequence 8, Application US/08978523
Patent No. 5883229
GENERAL INFORMATION:
APPLICANT: Guastella, John
TITLE OF INVENTION: Genes Coding For Bcl-y, a Bcl-2
TITLE OF INVENTION: Homologue
'NUMBER OF SEQUENCES: 53
CORRESPONDENCE ADDRESS:
ADDRESSE: STRRE, KESSLER, GOLDSTEIN & FOX P.L.L.C.
STREET: 1100 New York Avenue, N.W., Suite 600
                                                         780000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   480000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                    Conservative:
Mismatches:
Indels:
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Matches:
Conservative:
                                                                                                                                                             US-09-375-514-22 (1-205) x US-08-798-897-8 (1-24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches:
Indels:
                                                       Length:
Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US/08/978,523
FILING DATE: herewith
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/798,897
FILING DATE: February 11, 1997
CLASSIFICATION NUMBER: US 08/798,897
FILING DATE: February 11, 1997
CLASSIFICATION NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 32,893
REFERENCE/DOCKET NUMBER: 1483.01400(
TELEPHONE: 202-371-2600
TUBLESHONE: 202-371-2600
                                                                                                                                                                                              188 TrpileGlnAspAsnGlyGlyTrp 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        188 TrpileGlnAspAsnGlyGlyTrp 195
                                                                                                                                                                                                                 24 TGGATHCARGAYAAYGGNGGNTGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 202-371-2540
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                 55.00
100.00%
100.00%
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100.00%
100.00%
4.98%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: both
MOLECULE TYPE: CDNA
                                                                               Percent Similarity:
Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20005
                                Alignment Scores:
                                                                                                                                                                                                                                                                             US-08-978-523-8/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alignment Scores:
US-08-798-897-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-978-523-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match:
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                                                                                                                                                                                                                                                                RESULT 4
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APPLICANT: Reed, John
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES FOR INHIBITING THE
TITLE OF INVENTION: GROWTH OF CELLS EXPRESSING THE HUMAN BCL-2 GENE
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
ADDRESSEE: P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 4, Application US/08465485A
; Patent No. 5831066
; GENERAL INFORMATION:
APPLICANT: Reed, John
; TITLE OF INVENTION: Regulation of bcl-2 Gene Expression
; NUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARES PATENTIN RELEASE #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/217,082A
FILING DATE: 24-MAR-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/288,692
FILING DATE: 22-DEC-1988
ATTORNEY/AGENT INFORMATION:
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80000
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Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-375-514-22 (1-205) x US-08-217-082A-4 (1-33)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels:
                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMFUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                       Sequence 4, Application US/08217082A
Patent No. 5734033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Fortney, Andrew D.
REGISTRATION NUMBER: 34,600
REFERENCE/DOCKET NUMBER: 3335-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (408) 436-2075
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 33 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: P.C. STREET: 224 Airport Parkway CITY: San Jose STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.28e+03
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100.00%
4.26%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A. ZIP: 95110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Percent Similarity:
                    US-08-217-082A-4
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RESULT 5
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Sequence 4, Application US/09080285

Sequence 4, Application US/09080285

Fatent No. 6040181

APPLICANT: Reed, John

TITLE OF INVENTION: Regulation of bc1-2 Gene Expression

NUMBER OF SEQUENCES: 29

CORRESPONDENCE ADDRESS:

ADDRESSEE: DC.

STREET: 1755 S. Jefferson Davis Hwy., Suite 400

CITT: Alington

CONTROLL OF ADDRESSEE: VIC.

STREET: VIC.

CONTROLL OF ADDRESSEE: VIC.

CONTROLL OF ADDRESS
              ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, ADDRESSEE: P.C. STREET: 1755 S. Jefferson Davis Hwy., Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25 SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/465,485A FILING DATE: 05-JUN-1995 CLASSIFCATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-375-514-22 (1-205) x US-08-465-485A-4 (1-33)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3335-070-55 CONT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/288,692
FILING DATE: 22-DEC-1988
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/124,256
FILING DATE: 20-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 33:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (408) 436-2070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Fortney, Andrew D. REGISTRATION NUMBER: 34,600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA (genomic)
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.28e + 03
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100.00%
100.00%
4.26%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (408) 436-2075
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                   Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 33 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
CORRESPONDENCE, ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Virginia COUNTRY: U.S.A. ZIP: 22202
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                                                                                                                                      Arlington
                                                                                                                                                                                Virginia
                                                                                                                                                                                                              COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Alignment Scores:
                                                                                                                                                                                                                                            22202
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-465-485A-4
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                                                                                                                                              CITY: A
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Sequence 4, Application US/09724426
| Sequence 4, Application US/09724426
| Patent No. 6414134
| GENERAL INFORMATION:
| APPLICANT: Reed, John
| TILE OF INVENTION: Regulation of BCL-2 Gene Expression
| TILE OF INVENTION: Regulation of BCL-2 Gene Expression
| FILE REFERENCE: 10412-024
| CURRENT PFLIATOR DATE: 2000-11-28
| NUMBER OF SEQ ID NOS: 29
| SOFTWARE: PatentIn version 3.0
| SEQ ID NO 4
| LENGTH: 33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         333
   MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/080,285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-375-514-22 (1-205) x US-09-080-285-4 (1-33)
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps:
                                                                                                                                  CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/465,485
FILING DATE: 05-UNN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/124,256
FILING DATE: 20-SEP-1993
FILING DATE: 20-SEP-1993
FILING DATE: 21-FEB-1992
FILING DATE: 21-FEB-1992
PRIOR APPLICATION NUMBER: US 77/288,692
FILING DATE: 22-DEC-1988
FILING DATE: 22-DEC-1988
ATTORNEX/AGENT: NEVERMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       193 GlyGlyTrpValGlyAlaSerGly 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10 GGAGGCTGGGTAGGTGCATCTGGT 33
                                                                                                                                                                                                                                                                                                                                                                                       NAME: FOTTHRY, ANGTEW D.
REGISTRATION NUMBER: 34,600
REFERENCE/DOCKER NUMBER: 3335-
TELECOMMUNICATION INFORMATION:
TELEPRAX: (408) 436-2075
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 33 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.28e+03
47.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.28e+03
47.00
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100.00%
4.26%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acid
EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Alignment Scores:
                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alignment Scores:
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Sequence 8, Application US/07936421
| Patent No. 5750390
| GENERAL INFORMATION:
| APPLICANT: James D. Thompson APPLICANT: Kenneth G. Draper TITLE OF INVENTION: METHOD AND REAGENT FOR TITLE OF INVENTION: TREATMENT OF DISEASES CAUSED TITLE OF INVENTION: GENE NUMBER OF SEQUENCES: 22
| CORRESPONDENCE ADDRESS: ADDRESSE: Lyon & Lyon & Lyon STREET: 611 West SIxth Street
              Conservative:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/936,421
FILING DATE: 19920926
                             Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 19920826
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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Mismatches:
                                                                                     US-09-375-514-22 (1-205) x US-09-724-426-4 (1-33)
                                                  Indels:
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REPERENCE/DOCKET NUMBER: 197/243
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                      91 ProvalvalHisLeuAlaLeuArgGln 99
Percent Similarity: 100.008
Best Local Similarity: 100.008
Query Match: 4.268
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100.00%
4.08%
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Los Angeles
California
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Best Local Similarity:
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Sequence 11, Application US/08480994
Fatent No. 5834248
GENERAL INFORMATION:
APPLICANT: FALB, DEAN A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: FALB, DEAN A.
TITLE OF INVENTION: COMPOSITION AND METHODS FOR THE
TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
NUMBER OF SEQUENCES: 54
                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: THM PC compatible
OPERAITG SYSTEM: PC-DOS/MS-DOS
SUGTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/480,994
FILING DATE: 07-JUN-1995
CLASSIFICATION NUMBER: US 08/485,573
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485,573
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/485,573
FILING DATE: 07-JUN-1995
APPLICATION NUMBER: US 08/386,844
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Mismatches:
Indels:
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                                                                                                                                                                                            ADDRESSEE: PENNIE & EDMONDS
STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 08/386,844
FILING DATE: 10-FEB-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 11, Application US/08616844
Patent No. 5849578
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Coruzzi, Laura A. REGISTRATION NUMBER: 30,742
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 78
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (212) 869-8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
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(212) 869-8864
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Percent Similarity: 44.00
Best Local Similarity: 100.008
Query Match: 3.998
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                               New York
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US-08-480-994-11/c
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HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                   COUNTRY:
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LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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STATE:
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Sequence 11, Application US/08599654

Patent No. 5882925

GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
THE TRAINMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "synthetic oligonucleotide"
                                                                                                                                                                                                                                          COMPOTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPOTER: TBM FC Compatible
OPPERATING SYSTEM: FC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
FILING DATE: 15-MAR-1996
CLASSIFICATION: 800
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Matches:
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STREET: 1155 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSLYLCATION: OVO
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/599,654
FILING DATE: 09-FEB-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485,573
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/386,844
FILING DATE: 10-FEB-1995
ATTORNEY AGENT INFORMATION:
                       ADDRESSEE: PENNIE & EDMONDS STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7853-053
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100.00%
3.99%
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LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
CORRESPONDENCE ADDRESS:
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Best Local Similarity:
                                                                                                                                            New York
                                                                                                                                                                                                                   10036-2711
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                                                                                                         New York
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Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: Ne
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                                                                                                                                                                               COUNTRY:
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ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Patentin Release #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/599,654
FILING DATE: 09-FEB-1996
CLASSIFICATION: PANO
CLASSIFICATIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "synthetic oligonucleotide"
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Conservative:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
ADDRESSEE: PENNIE & EDMONDS
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 600

CLASSIFICATION: 600

PRIOR APPLICATION NUMBER: US 08/485,573

FILING DATE: 07-UN-1995

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/386,844

FILING DATE: 10-FEB-195

ATTORNEY/AGENT INFORMATION:
NAME: CORUZI, LAURA A

REGISTRATION NUMBER: 30,742

REPERENCE/COCKET NUMBER: 7853-041

TELEPHONE: (212) 790-9090

TELEPHONE: (212) 790-9090

TELEPAS: (212) 869-8864

TELEPAS: (212) 869-8864

TELERAY: OF SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: PID PC COMPATIBLE
COMPUTER: PID PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              140 AspGlyValAsnTrpGlyArg 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23 GATGGGGTGAACTGGGGGAGG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.54e+03
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Percent Similarity: 100.00%
Best Local Similarity: 100.00%
Query Match: 3.99%
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US-08-944-868A-11
                                                                                                                                                                                                                                                                        Alignment Scores:
                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                         Query Match:
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US-08-944-668A-11/C
Sequence 11, Application US/08944868A
Sequence 11, Application US/08944868A
Patent No. 6018025
TERENT INFORMATION:
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
CORRESPONDENCE ADDRESS:
ADDRESSEE: PENNIE & EDMONDS
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA

ZIP: 10036-2711

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMpatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFWMARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION NUMBER: US/08/944,868A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-375-514-22 (1-205) x US-08-485-573-11 (1-24)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length:
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION UNDRAR: US 08/386,844
FILING DATE: 10-FEB-1995
ATTORNEY/AGENT INFORMATION:
NAME: COTUZZI, LAUTA A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7853-032
TELECHOME: (212) 790-9090
TELECHAX: (212) 869-8864
TELEKAX: (6141 PENNIE
INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDENNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/386,844
FILING DATE: 10-FEB-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: CORUZZI, LAURA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7853-041
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/599,654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   140 AspGlyValAsnTrpGlyArg 146
                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23 GATGGGGTGAACTGGGGGAGG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                       .54e+03
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100.00%
3.99%
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                          Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                        5-08-485-573-11
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GENERAL INFORMATION:

APPLICANT: FALB, DEAN A
TILLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESSE: ADDRESSE: PENNIE & EDMONDS
STREET: 1155 Avenue of the Americas
                                                                                                                                                                           /desc = "synthetic oligonucleotide"
                                                                                                                                                                                                                                                                           4,0000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/944,423A
FILING DATE: 06-OCT-1997
                                                                                                                                                                                                                                                                                                               Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                 US-09-375-514-22 (1-205) x US-08-944-868A-11 (1-24)
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Matches:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/599, 654
FILING DATE: 09-FEB-1996
PRIOR APPLICATION NUMBER: US 08/485, 573
FILING DATE: JUN-07-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/485, 573
FILING APPLICATION DATA:
APPLICATION NUMBER: US 08/386, 844
                                                                                                                                                  MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "synthetic
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/386,844
FILING DATE: 10-FEB-1995
ATTORNEY_AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 11, Application US/08944423A Patent No. 6020463
                                                                                                                                                                                                                                                                                                                                                                                                                                        140 AspGlyValAsnTrpGlyArg 146
TELERAX: (212) 869-8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORUZZI, LAURA A
                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 78
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (212) 869-8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 11:
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100.00%
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3.99%
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(212) 869-8864
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Floppy disk
                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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ZIP: 10036-2711
COMPUTER READABLE FORM:
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TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER:
                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                           Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New York
New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 15
US-08-944-423A-11/C
                                                                                                                                                                                                                                                                                                       Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE:
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GENERAL INFORMATION:
APPLICANT: FALL, DEAN A
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
NUMBER OF SEQUENCES: 54
CORRESPONDENCE ADDRESS:
ADDRESSEE: PENNIE & EDMONDS LLP
STREET: 1155 Avenue of the Americas
CITY: New YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "synthetic oligonucleotide"
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            470000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/944,496
FILING DATE: 06-OCT-1997
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-375-514-22 (1-205) x US-08-944-496-11 (1-24)
                     US-09-375-514-22 (1-205) x US-08-925-743-11 (1-24)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 514

PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNBER: US 08/599,654

FILING DATE: 09-FEB-1996
PRIOR APPLICATION UNBER: US 08/485,573

APPLICATION UNBER: US 08/485,573

FILING DATE: 07-UN-1995

PRIOR APPLICATION UNBER: US 08/386,844

APPLICATION NUMBER: US 08/386,844

ATTIONE APPLICATION INFORMATION:
APPLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                             Sequence 11, Application US/08944496 Patent No. 6124433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      140 AspGlyValAsnTrpGlyArg 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: CORUZZI, LAURA A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 785
TELECOMMUNICATION INFORMATION:
                                                        140 AspGlyValAsnTrpGlyArg 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (212) 790-9090
TELERAX: (212) 869-8864
TELEX: 66141 PENIXE
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.54e+03
44.00
100.00%
100.00%
3.99%
                                                                          23 GATGGGGTGAACTGGGGGAGG 3
                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                         STATE: New York
                                                                                                                                              US-08-944-496-11/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match:
                                                                                                                                RESULT 17
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                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: FALB, DEAN A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7
7
0
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0
         420000
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COMPUTER: IBM PC compatible .
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/925,743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative:
Mismatches:
                                                                                                                                                                    Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                          US-09-375-514-22 (1-205) x US-08-944-423A-11 (1-24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches:
                                                                                                                                   Length:
Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SSEE: PENNIE & EDMONDS
7: 1155 Avenue of the Americas
New York
New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7853-032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/485,573
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               US-08-925-743-11/c
; Sequence 11, Application US/08925743
; Patent No. 6054558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: COCUZZI, LEURA A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7853
TELECOMMUNICATION INFORMATION:
TELEPAX: (212) 790-9090
TELERAX: (212) 869-8864
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                           DNA (genomic)
                                                                                                                              1.54e+03
44.00
100.00%
100.00%
3.99%
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100.00%
100.00%
3.99%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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Query Match:
                                                                                                                                                                  Percent Similarity:
Best Local Similarity:
Query Match:
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: lin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                   ; HYPOTHETICAL:
US-08-944-423A-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                      Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No.:
                                                                                                                                         Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                ESULT 16
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APPLICANT: Chenchik, Alex
Jokhadze, George
Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
EXPRESSION
NUMBER OF SEQUENCES: 1375
           METHOD OF ASSAYING DIFFERENTIAL EXPRESSION 1375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25
0
0
0
                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: Feat5EQ for Windows Version 2.0
SOFTWARE: Tast5EQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,998
FILING DATE: 21-MAY-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-375-514-22 (1-205) x US-08-859-998-596 (1-25)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: oligonucleotide primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length:
Matches:
     TITLE OF INVENTION: METHOD OF ASSAVING DI
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 1375
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION UNBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
TELECOMMUNICATION: INFORMATION:
TELEPHONE: 415-322-6770
TELEPHONE: 415-322-6775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 596, Application US/09225928 Patent No. 6352829 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  189 IleGlnAspAsnGlyGlyTrpVal 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24 ATCCAAGACCAGGGTGGTTGGGTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 415-854-0875
INPORMATION FOR SEQ ID NO: 596:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.45e+03
42.00
87.50%
87.50%
3.80%
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ZIP: 94025
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                         ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Menlo Park
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                           Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-225-928-596/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-859-998-596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Alignment Scores:
                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 20
                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: FALB, DEAN A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/925,767

FILING DATE: 09-SEPT-1997

CLASSIFICATION NUMBER: US 08/485,573

FILING DATE: 07-JUN-1995

RAPLICATION NUMBER: US 08/386,844

FILING DATE: 10-FEB-1995

APPLICATION NUMBER: US 08/386,844

FILING DATE: 10-FEB-1995

ATTORNEY/AGENT INFORMATION:

NAME: COLUZZI, LAURA A.

REGISTRATION NUMBER: 30,742

REFERENCE/DOCKET NUMBER: 7863-097

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-375-514-22 (1-205) x US-08-925-767-11 (1-24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length:
Matches:
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                                                                                                                                                                                                                                                                                                                             ADDRESSEE: PENNIE & EDMONDS
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 19
US-08-859-998-596/C
Sequence 596, Application US/08859998
Fatent No. 5994076
GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
APPLICANT: Jokhadze, George
APPLICANT: Bibliashvilli, Robert
                                             RESULT 18
US-08-925-767-11/c
; Sequence 11, Application US/08925767
; Patent No. 6255084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           140 AspGlyValAsnTrpGlyArg 146
23 GATGGGGTGAACTGGGGGAGG 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.54e+03
44.00
100.00%
100.00%
3.99%
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
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HYPOTHETICAL: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Percent Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S-08-925-767-11
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DB:
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GENERAL INFORMATION:
APPLICANT: CHITTENDEN, Thomas D.; and
APPLICANT: LUTZ, Robert J.
TITLE OF INVENTION: NOVEL PEPTIDES AND COMPOSITIONS WHICH
TITLE OF INVENTION: MODULATE APOPTOSIS
NUMBER OF SEQUENCES: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; SEQUENCE DESCRIPTION: Oligonucleotide primer; SEQUENCE DESCRIPTION: SEQ ID NO: 596: US-09-225-928-596
    COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/225,928
FILING DATE: 05-Jan-1999
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/440,391
FILING DATE: 12-MAY-1995
                                                                                                                                                                     APPLICATION NUMBER: 08/659,998
FILING DATE: 21-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
TELEPHONE: 415-322-5070
TELEPHONE: 415-322-5070
TELERAX: 415-884-0875
INFORMATION FOR SEQ ID NO: 596:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-375-514-22 (1-205) x US-09-225-928-596 (1-25)
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Hale and Dorr
STREET: 1455 Pennsylvania Avenue, N.W.
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                104322.147
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24 ATCCAAGACCAGGGTGGTTGGGTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: WIXON, HENRY N.
REGISTRATION NUMBER: 32,073
REFERENCE/DOCKET NUMBER: 10.
                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 25 base pairs
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.45e+03
42.00
87.50%
87.50%
3.80%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SIATE: D.C.
STATE: D.C.
20004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-440-391-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE
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US-08-908-597A-21

Sequence 21, Application US/08908597A

Sequence 21, Application US/08908597A

Patent No. 5863795

GENERAL INFORMATION:
APPLICANT: CHITTENDEN, Thomas D.; and
APPLICANT: CHITTENDEN, Thomas D.; and
APPLICANT: LUTZ, Robert J.

TILLE OF INVENTION: NOVEL PEPTIDES AND COMPOSITIONS WHICH
TITLE OF INVENTION: MODULATE APOPTOSIS
NUMBER OF SEQUENCES: 34

CORRESPONDENCE ADDRESS:

CORRESPONDENCE ADDRESS:

STREET: 1455 Pennsylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       39
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APPLICATION NUMBER: US/08/908,597A
                                                                                                                                                                                                                                                                                                        Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   101 GlyAspAspPheSerArgArgTyrArgGlyAspPhe 112
                                                                                                                                                                                                                                                                                                                                                                                                            US-09-375-514-22 (1-205) x US-08-440-391-21 (1-39)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION: 530
PRIOR APPLICATION DATA:
PILING DATE: 12-MAY-1995
ATTORNEY INFORMATION:
NAME: WIXON, HENRY N.
REGISTRATION NUMBER: 32,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   104322.147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-942-8400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-942-6400
TELEPHONE: 202-942-8484
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 1455 Pennsylvania CITY: Washington STATE: D.C. ZTATE: D.C. ZIP: 20004 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                               ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-440-391-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                     .52e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 202-942-8484
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                      42.00
75.00%
58.33%
3.80%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 39 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                      LENGTH: 39 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                      Percent Similarity:
Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-908-597A-21
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                                                                                                                                                                                                                                                      Alignment Scores:
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                                                                                                                                                                                                                                                                         Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 22
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TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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Query Match:
                                                                                                                                                                                                                                                                    FILING DATE: H
                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: 11:
MOLECULE TYPE:
                                                                                                 STATE: D.C. 20004
                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCT-US96-06122-21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-426-819A-28
                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (C) ATTORNEY DOCKET NO. 104322.147CIP TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                        LUTZ, Robert J.
TITLE OF INVENTION: NOVEL PEPTIDES AND COMPOSITIONS WHICH MODULATE APOPTOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 20004

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA: 25-7an-1999

FILING DATE: 25-7an-1999

CLASSIFICATION: CONKNOWN>

ATTORNEY/ABGENT INFORMATION:

NAME: WIXON, HENRY N.

REGISTRATION NUMBER: 32,073

REGISTRATION NUMBER: 32,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: IMMUNOGEN, INC.
TITLE OF INVENTION: NOVEL PEPTIDES AND COMPOSITIONS
        Conservative:
                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Hale and Dorr
STREET: 1455 Pennsylvania Avenue, N.W.
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative:
Mismatches:
                                                                                US-09-375-514-22 (1-205) x US-08-908-597A-21 (1-39)
                       Mismatches:
Indels:
                                                                                                                    US-09-375-514-22 (1-205) x US-09-236-385A-21 (1-39)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              101 GlyAspAspPheSerArgArgTyrArgGlyAspPhe 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches:
                                                                                                                                                                                                                                                                         APPLICANT: CHITTENDEN, Thomas D.; and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID NO: 21:
                                                                                                                                                                                                                       Sequence 21, Application US/09236385A Patent No. 6221615 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21, Application PC/TUS9606122 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 39 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 202-942-84(
TELEFAX: 202-942-8484
INFORMATION FOR SCO ID NO: 21:
SEQUENCE CHARACTERISTICS:
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   75.00%
58.33%
3.80%
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75.00%
58.33%
3.80%
                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: D.C.
Percent Similarity:
Best Local Similarity:
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Query Match:
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                                     Query Match:
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Sequence 28, Application US/08426819A
Patent No. 5723318
GENERAL INFORMATION
APPLICANT: Vojima, Tetsuo
APPLICANT: Kojima, Tetsuo
APPLICANT: Nojima, Tetsuo
APPLICANT: Nojima, Tetsuo
TITLE OF INVENTION: Genes Coding for Megakaryocyte
TITLE OF INVENTION: Potentiator
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                              OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/06122
FILING DATE: HEREWITH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative:
Mismatches:
WHICH MODULATE APOPTOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-375-514-22 (1-205) x PCT-US96-06122-21 (1-39)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSE: Birch, Stewart, Kolasch & Birch STREET: P.O. Box 747
CITY: Falls Church STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  104322.147PCT
                                                         E: Hale and Dorr
1455 Pennsylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels:
                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/440,391
                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: WIXON, HENRY N.
REGIGSTRATION NUMBER: 32,073
REFERENCE/DOCKET NUMBER: 1043
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-942-8400
TELEFAX: 202-942-8484
                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 22040-0747
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 12-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.52e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  42.00
75.00%
58.33%
3.80%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 39 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
               NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                      COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                Washington
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TITLE OF INVENTION: Metal Resistance Sequences and TITLE OF INVENTION: Transgenic Plants
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Greenlee, Winner and Sullivan, P.C. STREET: 5370 Manhattan Circle, Suite 201 CITY: Boulder
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US08/426,819A
                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "synthetic primer, 850GS"
                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-375-514-22 (1-205) x US-08-426-819A-28 (1-33)
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21-APR-1995
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FILING DATE: 21-APR-1995
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                                                                                              ATTORNEY AGENT INFORMATION:
NAME: MUTPHY Jr., Gerald M.
REGISTRATION NUMBER: 28,971
REPERENCE/DOCKET NUMBER: 230-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-205-8000
TELEFRAX: 703-205-8000
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Meagher, Richard B. APPLICANT: Sommers, Anne O.
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                                                                                                                                                                                                                                                                                                                                                                                                                            41.00
80.00%
80.00%
3.71%
                                                                                                                                                                                                                                                        LENGTH: 33 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                     CLASSIFICATION:
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                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                Percent Similarity:
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SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: PSPLICATION NUMBER: US/08/878,957 FILING DATE: 19-JUN-1997 CLASSIFICATION: 800 PRIOR APPLICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Meagher, Richard B.
APPLICANT: Summers, Anne O.
APPLICANT: Stander Sudy, Clayton L.
TITLE OF INVENTION: Metal Resistance Sequences and TITLE OF INVENTION: Transgenic Plants
NUMBER OF SEQUENCES: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Greenlee, Winner and Sullivan, P.C. 5370 Manhattan Circle, Suite 201
                                                                                                                                                                                               Conservative:
                                                                                                                                                                                                                    Mismatches:
                                                                                                                                                                                                                                                                                       US-09-375-514-22 (1-205) x US-08-427-097-6 (1-36)
                                                                                                                                                                Length:
Matches:
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DESCRIPTION: /desc = "Olgionucleotide"
                                                                                                                                                                                                                                     Indels:
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Olgionucleotide"
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                        Gabs:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/427,097
FILING DATE: 21-APR-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                            75 ProAlaAlaProGlyAlaAlaAlaGly 83
                                                                                                                                                                                                                                                                                                                                                 35 CCIGCIGCCGGGIGAGGCIGCAGGA 9
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 6, Application US/08878957
Patent No. 5965796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Ferber, Donna M. REGISTRATION NUMBER: 33,878 REFERENCE/DOCKET NUMBER: 40-TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (303) 499-8080
TELEFAX: (303) 499-8089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.97e+03
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LENGTH: 36 base pairs
                                                                                                                                                                                   41.00
88.89%
88.89%
3.71%
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CITY: Boulder
STATE: Colorado
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STRANDEDNESS:
                                                                                                                                                                                                        Percent Similarity:
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Pred. No.:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-878-957-6/c
                                                                                                                                                      Alignment Scores:
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                                                                                                               US-08-427-097-6
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us-09-375-514-22.rni

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44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro 59
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/145,704
FILING DATE: 28-OCT-1993
CLASSIFICATION NUMBER: US/08/145,704
PRIOR APPLICATION S14
PRIOR APPLICATION NUMBER: US/08/053,027
FILING DATE: 23-APR-1993
ATTONNEY/AGERT INFORMATION:
NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
REFERENCE/POCKET NUMBER: 32,714
RELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/651-5151
merrey. 713/651-5246
        80400
                    Conservative:
Mismatches:
                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Rando, Robert F.
APPLICANT: Rennewald, Susan
APPLICANT: Zendegul, Joseph G.
APPLICANT: Joshua O. Ojwang
TITLE OF INVENTION: Anti-Viral Guanosine-Rich
TITLE OF INVENTION: Oligonucleotides
NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative:
Mismatches:
                                                                                         US-09-375-514-22 (1-205) x US-08-878-957-6 (1-36)
   Matches:
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                                                  Indels:
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Matches:
                                                                     Gaps:
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                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Fulbright & Jaworski
STREET: 1301 McKinney, Suite 5100
                                                                                                                         75 ProAlaAlaProGlyAlaAlaAlaGly 83
                                                                                                                                            35 CCTGCTGCGCGGGTGAGGCTGCAGGA 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                    RESULT 28
US-08-145-704-2/c
; Sequence 2, Application US/08145704
; Patent No. 5567604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                41.00
88.89
88.89
3.71
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50.00%
50.00%
3.71%
           Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U.S.A.
ZIP: 77010-3095
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                   Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Texas
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                                               Query Match:
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44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro 59
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APPLICANT: Rando, Robert F.
APPLICANT: Fennewald, Susan
APPLICANT: Fennewald, Susan
APPLICANT: Ojwang, Joseph G.
APPLICANT: Hogan, Michael E.
TITLE OF INVENTION: Anti-Viral Guanosine-Rich
ITTLE OF INVENTION: Ojigonucleotides
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER REALABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/987,574
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/04529
FILING DATE: 28-OCT-1993
APPLICATION NUMBER: US 08/053,027
FILING DATE: 23-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 32,714
REFERENCE/POCKET UNMBER: D-5574-CIP
TELECOMMUNICATION:
TELEPHONE: 713,651-5151
                                                                                                                                                                                                                                                                       ADDRESSEE: Fulbright & Jaworski
STREET: 1301 McKinney, Suite 5100
RESULT 29
US-08-987-574-2/C
; Sequence 2, Application US/08987574
; Patent No. 6150339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2, Application US/08535168; Patent No. 6184369; GENERL INFORMATION: APPLICANT: Rando, Robert F. APPLICANT: Fennewald, Susan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  41.00
50.00%
50.00%
3.71%
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TELEX: 762829
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                     CITY TEXAL COUNTRY: U.S.A. 77010-3095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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STREET: 150.
THV: Houston
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match:
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600 Travis, Suite 1800

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44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro 59
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APPLICANT: Rando, Robert F.
APPLICANT: Ojwan, Michael E.
APPLICANT: Wallace, Thomas L.
APPLICANT: Wallace, Thomas L.
APPLICANT: Wallace, Thomas L.
TITLE OF INVENTION: Anti-Viral Guanosine-Rich
TITLE OF INVENTION: Tetrad Forming Oligonucleotides
NOMBER OF SEQUENCES:
ADDRESSEE: Conley, Rose & Tayon, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTER TO S.A.

ZIP: 77010-3095

ZIP: 77010-3095

COMPUTER REDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-POS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION NUMBER: US/08/535,168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length:
Matches:
Conservative:
APPLICANT: Zendegui, Joseph G.
APPLICANT: Ojwang, Joshua O.
APPLICANT: Hogan, Michael E.
TITLE OF INVENTION: Anti-Viral Guanosine-Rich
TITLE OF INVENTION: Oligonucleotides
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fulbright & Jaworski
STREET: 1301 MCKInney, Suite 5100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-375-514-22 (1-205) x US-08-535-168-2 (1-38)
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                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFTCATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 97/US94/04529
FILING DATE: 28-0CT-1993
APPLICATION NUMBER: 08 08/053,027
FILING DATE: 23-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: PAUL, FIDMES 13-714
REGISTRATION NUMBER: 32-714
REFERENCE/DOCKET NUMBER: D-5574-CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 2, Application US/09017974 Patent No. 6288042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/651-5151
TELEFAX: 713/651-5246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 713/651-5246
TELEX: 762829
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 38 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5.36e+03
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3.71%
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Best Local Similarity:
                                                                                                                                                                        STREET: LOUE ..
                                                                                                                                                                                                                          Texas
                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-017-974-2/c
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Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-535-168-2
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44 ProAlaProGly1lePheSerSerGlnProGlyHisThrProHisPro 59
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APPLICANT: Fennewald, Susan
APPLICANT: Pennewald, Susan
APPLICANT: Olwang, Joshua O.
APPLICANT: Olwang, Joshua O.
APPLICANT: Hogan, Michael E.
APPLICANT: Pommier, Eyves
APPLICANT: Pommier, Eyves
APPLICANT: Mazumder, Abhijit
TITLE OF INVENTION: Anti-Viral Guanosine-Rich
TITLE OF INVENTION: Oligonucleotides
NUMBER OF SEQUENCES: 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-375-514-22 (1-205) x US-09-017-974-2 (1-38)
                                                                                                                      OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS Word 97 (saved as .txt file)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length:
                                                                                                                                                                                                                                                                                                                    RILING DATE: 09-DEC-97
ATTORNEY/AGENT INFORMATION:
NAME: McDaniel, C. Steven
REGISTRATION NUMBER: 33,962
REFERENCE/DOCKET NUMBER: 1472-06223
REFERENCE/DOCKET NUMBER: 1472-06223
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/238-8018
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Conley, Rose & Tayon, P.C. STREET: 600 Travis, Suite 1850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps:
                                                                                                                                                                                      APPLICATION NUMBER: US/09/017,974
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; Patent No. 6323185
                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/037,374
FILING DATE: 04-FEB-97
APPLICATION NUMBER:
                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5.36e+03
41.00
50.00%
50.00%
3.71%
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LENGTH: 38 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                    : Texas
RY: U.S.A.
77002-2912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Texas
Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Percent Similarity:
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US-08-682-255A-2/C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-017-974-2
                                       COUNTRY:
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                    STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match:
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44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro 59
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Matches:
Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Anti-viral Guanosine-Rich Oligonucleotides
COMPUTER: IBM PC compatible OPERATING SYSTEM: MS Windows 95 SOFTWARE: MS WORD 97 (saved as .txt file) CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/682,255A FILING DATE: 17-JULY-1996 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-375-514-22 (1-205) x US-08-682-255A-2 (1-38)
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ADDRESSEE: Conley, Rose & Tayon, P.C.
STREET: 600 Travis, Suite 1850
CITY: Houston
STATE: Texas
                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/535,168
FILING DATE: 23-007-95
APPLICATION NUMBER: 60/001,505
FILING DATE: 19-JULY-95
APPLICATION NUMBER: 60/014,007
FILING DATE: 25-MARCH-96
APPLICATION NUMBER: 60/013,688
FILING DATE: 19-MARCH-96
APPLICATION NUMBER: 60/015,714
FILING DATE: 17-APRIL-96
APPLICATION NUMBER: 60/015,714
FILING DATE: 23-APRIL-96
APPLICATION NUMBER: 60/015,714
FILING DATE: 23-APRIL-96
APPLICATION NUMBER: 60/015,714
FILING DATE: 23-APRIL-96
APPLICATION NUMBER: 100/16,271
FILING DATE: 12-APRIL-96
APPLICATION NUMBER: 100/16,271
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Zendegui, Joseph G.
Ojwang, Joshua O.
Hogan, Michael E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/09429130
Patent No. 6355785
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pommier, Eyves
Mazumder, Abhijit
60/015,714
                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: MCDaniel, C. Steven
REGISTRATION NUMBER: 33,962
REFERENCE/DOCKET NUMBER: 147
TELECOMMUNICATION:
TELEPHONE: 713/238 8010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Rando, Robert F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 713/238-8008
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5.36e+03
41.00
50.00%
50.00%
3.71%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
LENGTH: 38 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: 713/238-8008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A. ZIP: 77002-2912
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
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PCT-US92-09202-10/c

Sequence 10, Application PC/TUS9209202

GENERAL INFORMATION:

APPLICANT: Hogan, Michael E.

TITLE OF INVENTION: Triplex Forming Oligonucleotide Reagents Targeted

TITLE OF INVENTION: to the Neu Oncogene Promoter and Method of Use

NUMBER OF SEQUENCES: 10

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: MS WORD 97 (saved as .txt file)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: MCDaniel, C. Steven
REGISTRATION NUMBER: 33,962
REFERENCE/COCKET NUMBER: 1472-06214
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/238-8010
TELEFAX: 713/238-8008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-375-514-22 (1-205) x US-09-429-130-2 (1-38)
                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/429,130
FILING DATE: 28-Oct-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                           Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-09-429-130-2
                                                                                                                                                                                                                                                 23-APRIL-96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Fulbright & Jaworski
STREET: 1301 McKinney, Suite 5100
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 38 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTEX: U.S.A.
ZIP: 77010-3095
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   41.00
50.00%
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3.71%
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Best Local Similarity:
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APPLICANT: Tsai, Ming-Jer
APPLICANT: Hogan, Michael H
APPLICANT: Hogan, Michael H
APPLICANT: O'Malley, Bert W
APPLICANT: Ing, Nancy H
TITLE OF INVENTION: Novel Triplex Forming Oligonucleotides
TITLE OF INVENTION: and Methods for their use
CORRESPONDENCE: 18
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            44 ProAlaProGly1lePheSerSerGlnProGlyHisThrProHisPro 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Fulbright & Jaworski Patent Department
1301 McKinney #5100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08329
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/09202
FILING DATE: 19921028
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative:
                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: Triplex forming oligonucleotide HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-375-514-22 (1-205) x PCT-US92-09202-10 (1-38)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Adler, Benjamin A
REGISTRATION NUMBER: 35,423
REFERENCE/DOCKET NUMBER: d5486
TELEPOMUNICATION INFORMATION:
TELEPHONE: 713-651-5151
TELEPRAX: 713-651-5346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCT-US93-08329-11/c
; Sequence 11, Application PC/TUS9308329
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                           ATTORNEY AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
REFERENCE/DOCKET NUMBER: D-5393
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/651-5151
                                                                                                                                                                                                                                              TELEX: 713/v--
TELEX: 762839
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 38 base pairs
"YPE: NUCLEIC ACID
"YPE: NUCLEIC ACID
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50.00%
50.00%
3.71%
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TELEPHONE: 713/651-5246
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                                                                                                                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 77010-3095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 1301 m
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
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PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/535,168; 60/001,505; 60/014,007; 60/013,688; APPLICATION NUMBER: 60/015,714; 60/016,271

FILING DATE: 23-0CT-95; 17-JULY-96; 25-MARCH-96; 19-MARCH-96; 23-FILING DATE: APRIL-96; 17-APRIL-96

ATTORNEY APPLICATION:
                                                                                                                                                                                                                                                                                                                                                                                            44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro 59
                                                                                                                                                                                                                                                                                                                                                                                                                1620838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application PC/TUS9611786

GENERAL INFORMATION:
APPLICANT: Rendo, Robert F.
APPLICANT: Fennewald, Susan
APPLICANT: Zendegui, Joseph G.
APPLICANT: Powmier, Boshua O.
APPLICANT: Powmier, Eyves
APPLICANT: Powmier, Eyves
APPLICANT: Maxumder, Abhijit
TITLE OF INVENTION: Anti-Viral Guanosine-Rich
TITLE OF INVENTION: Oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/11786
FILING DATE: 17-UULY-1996
                                                                                                                                                                                                                                            Matches:
Conservative:
                                                                                                                                                                                                                                                                                                                                                        US-09-375-514-22 (1-205) x PCT-US93-08329-11 (1-38)
                                                                                                                                                                                                                                                                                   Mismatches:
                                                                                                                                                                                                                                                                                                     Indels:
                                                                                                                                                                                                                          Length:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: McDaniel, C. Steven
REGISTRATION NUMBER: 33,962
REFERENCE/DOCKET NUMBER: 1472-06214
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                           TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Floppy disk
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50.00%
50.00%
3.71%
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                    SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
INFORMATION FOR SEQ ID NO:
                                    LENGTH: 38 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                          single
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Best Local Similarity:
Query Match:
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STREET: 600
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                                                                    STRANDEDNESS:
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PCT-US93-08329-11
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linear

TOPOLOGY:

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CUBA
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                                                                                                                                                                                                                              CITY:
STATE:
                                                                                                                                                                                                                                                                      PCT-US95-06987-10/C
Sequence 10, Application PC/TUS9506987
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Santon, June
TITLE OF INVENTION: The Inhibition of Cytokine Production by TITLE OF INVENTION: Oligonuclectides and Inhibition of Tumors Which are NUMBER OF SEQUENCES: 10
                                                                                                                                                                                                44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro 59
                                                                                                                                                                                                                    27
0
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/06987
                                                                                                 Conservative:
Mismatches:
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Conservative:
                                                                                                                                                                   US-09-375-514-22 (1-205) x PCT-US96-11786-2 (1-38)
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                                                                  Length:
Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Fulbright & Jaworski
STREET: 1301 McKinney, Suite 5100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
REFERENCE/DOCKET NUMBER: D-5669
TELECOMOUNICATION INFORMATION:
TELEPHONE: 713/651-5151
TELEFAX: 713/651-5246
    DNA (genomic)
                                                            5.36e+03
41.00
50.00%
50.00%
3.71%
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40.00
85.71%
85.71%
3.62%
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                         Percent Similarity:
Best Local Similarity:
Query Match:
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77010-3095
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Houston
; MOLECULE TYPE:
PCT-US96-11786-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Texas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                               Alignment Scores:
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TELEX: 7
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TITLE OF INVENTION: Method for obtaining modified
TITLE OF INVENTION: Immunoglobulins with reduced immunogenicity of murine
TITLE OF INVENTION: antibody variable domains, compositions containing them.
NUMBER OF SEQUENCES: 31
CORRESPONDENCE ADDRESS:
ADDRESSEE: CENTRO DE INVINOLOGIA MOLECULAR
STREET: 215 Y 15, ATABEY PLAYA
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TITLE OF INVENTION: immunoglobulins with reduced immunogenicity of murine
TITLE OF INVENTION: antibody variable domains, compositions containing them.
NUMBER OF SEQUENCES: 31
                                                                                                                                                                                                                                                                                                                                        COUNTIEST CUDAR
CIDINITATION
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PAPPLICATION DATA:
APPLICATION NUMBER: US/08/497,312
FILING DATE: 30-JUN-1995
APPLICATION NUMBER: CU 80/94
FILING DATE: 30-JUN-1994
ATTORNEY/AGENT INFORMATION:
NAME: BOND, LAURENCE B.
REDISTRATION NUMBER: 30,549
FREISTRATION NUMBER: 30,549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                35
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Mismatches:
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ADDRESSEE: CENTRO DE INMUNOLOGIA MOLECULAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-375-514-22 (1-205) x US-08-497-312-8 (1-35)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       45 AlaProGlyIlePheSerSerGlnProGly 54
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US-08-497-312-8

Sequence 8, Application US/08497312

Patent No. 5712120

GENERAL INFORMATION:

APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-497-312-10/c
; Sequence 10, Application US/08497312
; Patent No. 5712120
; GENERAL INFORMATION:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 26.
TELECOMUNICATION INFORMATION:
TELEPHONE: 801/532-1922
TELEFAX: 801/531-9168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               40.00
80.00%
70.00%
3.62%
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 35 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: unkno
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Best Local Similarity:
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US-09-375-514-22 (1-205) x US-08-469-318-110 (1-40)
        ATTORNEY AGENT INFORMATION:
NAME: White, John P. 28,678
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 36537-B2-Y
TELECOMUNICATION INFORMATION:
TELEPHONE: (212) 278-0400
TELEPHONE: (212) 391-0525
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "synthetic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             % Sequence 110, Application US/08469318 Sequence 110, Application US/08469318 Fatent No. 6022535 GENERAL INFORMATION:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/446,872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
INFORMATION FOR SEQ ID NO: 110:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         53 ProGlyHisThrProHisPro 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 CCAGGACGAACGCCACATCCA 21
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40.00
85.718
85.718
3.628
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100.00%
75.00%
3.53%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
CLASSIFICATION: 514
                                                                                                                                                                                                                                 nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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Best Local Similarity:
Query Match:
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Query Match:
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                                                                                                                                                                                                                                 TYPE: nucleic STRANDEDNESS:
                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Percent Similarity:
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US-08-463-682-7
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Patent No. 6008193
GENERAL INFORMATION:
GAPPLICANT: Leonard Garfinkel, et al.
TITLE OF INVENTION: Cloning and Production of Human Von
TITLE OF INVENTION: Willebrand Factor GPID Binding Domain Polypeptides and
TITLE OF INVENTION: Methods of Using Same
                                                                                                                                                                                     CURRENT APPLICATION DATA:

SOFTWARE: Petentin Release #1.0, Version #1.30 (EPO)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/497,312

FILING DATE: 30-UDN-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/44

ATTORNEY/AGENT INFORMATION:

NAME: BOND, LAURENCE B.

REGISTRATION NUMBER: 30,549

REFERENCE/DOCKET NUMBER: 2629US

TELECOMMUNICATION INFORMATION:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPPRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
FILING DATE: 05-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches:
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            32 GCCCCAGGCTTCTTCACTTCAGCCCCAGGC 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Cooper & Dunham
1185 Avenue of Americas
                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
      215 Y 15, ATABEY PLAYA
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40.00
80.00%
70.00%
3.62%
                                                                                                                                  Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 36 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 2.
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: New York COUNTRY: U.S.A. ZIP: 10036 COMPUTER READABLE FORM:
                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity:
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MOLECULE TYPE: CI
HYPOTHETICAL: NO
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                                                                      COUNTRY: CUBA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                            HAVANA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Percent Similarity:
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TITLE OF INVENTION: Multivariant IL-3 Hematopoiesis Fusion TITLE OF INVENTION: Protein NUMBER OF SEQUENCES: 196
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO) CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/469,318
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0 0 0 0 0 0
Length:
Matches:
Conservative:
Mismatches:
Indels:
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Mismatches:
                                                                                            US-09-375-514-22 (1-205) x US-08-463-682-7 (1-40)
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APPLICANT: Brader, S. C. APPLICANT: Braford-Goldberg, Sarah R. APPLICANT: Caparon, Malre H. APPLICANT: Caparon, Malre H. APPLICANT: Caparon, Alan M. APPLICANT: Klein, Barbara K. APPLICANT: Klein, Barbara K. APPLICANT: Nickarn, John P. APPLICANT: Paik, Kumnan APPLICANT: Thomas, John W. TITLE OF INVENTION: Multivariant IL-3 Hematopoiesis TITLE OF INVENTION: Fusion Protein NUMBER OF SEQUENCES: 197
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,872A
FILING DATE: 06-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION 1995
FILING DATE: 14-FEB-1994
ATPOREY/AGENT INFORMATION:
NAME: Bennett, Dennis A.
REGISTRATION NUMBER: 34,547
REFERENCE/DOCKET NUMBER: 34,547
REFERENCE/DOCKET NUMBER: C-2790/1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Dennis A. Bennett, G.D. Searle & Co., ADDRESSEE: Corporate Patent Dept.
STREET: P. O. Box 5110
CITY: Chicago
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Mismatches:
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DESCRIPTION: /desc = "synthetic DNA"
                           Sequence 110, Application US/08446872A; Patent No. 6361977; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-762-227A-110/c
; Sequence 110, Application US/08762227A
; Patent No. 6436387
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (314)737-6986
TELEFAX: (314)737-6972
INFORMATION FOR SEQ ID NO: 110:
                                                                                        Abrams, Mark A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8.67e+03
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100.00%
75.00%
3.53%
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LENGTH: 40 base pairs
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STRANDEDNESS: single
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STATE: Illinois
COUNTRY: USA
ZIP: 60680
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Best Local Similarity:
          JS-08-446-872A-110/C
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Thomas, John W. TITLE OF INVENTION: Fusion Proteins Comprising Multiply Mutated Inteleukin-3 (IL-CORRESPONDENCES: 197
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Dennis A. Bennett, G.D. Searle & Co., ADDRESSEE: Corporate Patent Dept.
STREET: P. O. Box 5110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPOURE READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
CSOTWARE: PATENTIN Release #1.0, Version #1.25
APPLICATION NUMBER: US/08/468,609A
FILING DATE: 06-JUN-1995
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Matches:
Conservative:
Mismatches:
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                                                                                                                                                                                             Bauer, S. C.
Braford-Goldberg, Sarah R.
Caparon, Maire H.
Easton, Alan M.
Klein, Barbara K.
McKearn, John P.
Olins, Peter O.
Paik, Kumnan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/192,325
FILING DATE: 14-FEB-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Bennett, Dennis A.
REGISTRATION NUMBER: 34,547
REFERENCE/DOCKET NUMBER: C-2790/3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (314)/37-6986
TELEFAX: (314)/737-6986
                                                                                                                 Sequence 110, Application US/08468609A Patent No. 6030812 GENERAL INFORMATION:
84 ProAlaLeuSerProValProPro 91
                     38 CCTGCCATGGCTCCAGTACCACCA 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 110: SEQUENCE CHARACTERISTICS: LENGTH: 40 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                  Abrams, Mark A.
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100.00%
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3.53%
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CITY: Chicago
STATE: Illinois
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Best Local Similarity:
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                                                                                              JS-08-468-609A-110/c
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APPLICANT:
APPLICANT:
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APPLICANT:
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TYPE: nucleic acid
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                                                                                                                                                                                                                                                                    ADDRESSEE: Dennis A. Bennett, G.D. Searle & Co., Corporate Patent Dept.
STREET: P. O. Box 5110
                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                Thomas, John W. TITLE OF INVENTION: Multivariant IL-3 Hematopolesis
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Mismatches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/762,227A
FILING DATE: 09-Dec-1996
CLASSIFICATION - CURROWN-
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/192,325
FILING DATE: 14-FEB-1994
APPLICATION NUMBER: US 08/446,872
FILING DATE: 06-JUN-1995
ATORNEY/AGENT INFORMATION:
NAME: Bennett, Dennis A.
REGISTRATION NUMBER: 34,547
REFERENCE/DOCKET NUMBER: C-2790/5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DESCRIPTION: /desc = "synthetic DNA" SEQUENCE DESCRIPTION: SEQ ID NO: 110:
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Matches:
                               Bauer, S. C.
Braford-Goldberg, Sarah R.
Caparon, Maire H.
Easton, Alan M.
Klein, Barbara K.
McKearn, John P.
Olins, Peter O.
Paik, Kumnan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels:
                                                                                                                                                                                                              NUMBER OF SEQUENCES: 197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 110, Application PC/TUS9501185 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: (708)470-6501
TELEFAX: (708)470-6881
INFORMATION FOR SEQ ID NO: 110:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           84 ProAlaLeuSerProValProPro 91
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
               APPLICANT: Abrams, Mark A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8.67e+03
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100.00%
75.00%
3.53%
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                                                                                                                                                                                                                                                                                                                                 CITY: Chicago
STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 60680
GENERAL INFORMATION:
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Best Local Similarity:
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APPLICANT: John Edward No. 6316196ris MORTEN
TITLE OF INVENTION: METHODS FOR ANALYZING LTC4 SYNTHASE POLYMORPHISMS
AND DIAGNOSTIC USE
TITLE OF INVENTION: Multivariant IL-3 Hematopoiesis Fusion TITLE OF INVENTION: Protein NUMBER OF SEQUENCES: 196
COMPTER READABLE FORM:
                                                               SOFTWARE: PAtentin Release #1.0, Version #1.30 (EPO) CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/0110F
FILING DATE: 02-PFR-100C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             60000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-375-514-22 (1-205) x PCT-US95-01185-110 (1-40)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSER: P111sbury Winthrop, L.L.P.
STREET: 1100 New York Ave., N.W.
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB98/02468
FILING DATE: Adgust 18, 1998
APPLICATION NUMBER: GB 9717766.1
FILING DATE: Adgust 22, 1997
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/485,636
FILING DATE: 15-Feb-2000
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: MS WORD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "synthetic DNA"
                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/192325
FILING DATE: 14-FEB-1994
INFORMATION FOR SEQ 1D NO: 110:
SEQUENCE CHARACTERISTICS:
LENGTH: 40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 2, Application US/09485636; Patent No. 6316196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
ZIP: 20005
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8.67e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               39.00
100.00%
75.00%
3.53%
                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                         PCT-US95-01185-110
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55 HisThrProHisProAlaAlaSer 62
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GENERAL INFORMATION:
APPLICANT: Rando, Robert F.
APPLICANT: Santon, June
TITLE OF INVENTION: The Inhibition of Cytokine Production by
TITLE OF INVENTION: Oligonucleotides and Inhibition of Tumors Which are
TITLE OF INVENTION: Responsive to Cytokine Factors
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             00700
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/06987
                                                                                                                                                                                                                                                                                                                                                                               Conservative:
Mismatches:
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Matches:
Conservative:
Mismatches:
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                                                                                                                                                                                                                                                                                                     Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels:
;
TOPOLOGY: linear

MOLECULE TYPE: other nucleic acid

SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-09-485-636-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Houston STATE: Texture 5100 STATE: Textu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   38 AlaProProGlyAlaAlaProAla 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29 GCCCGCCTGGAGCGCCCCCGGCC 6
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CLASSIFICATION:
CLASSIFICATION:
ATTORNEY AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTATION NUMBER: 32,714
REFERENCE/DOCKET NUMBER: D-56
TELECOMMUNICATION INFORMATION:
TELECHONE: 713,651-5151
TELEFRX: 713,651-5146
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TELERAX: 762829
INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS: LENGTH: 35 base pairs
"YPE: nucleic acid
"YPE: nucleic acid
                                                                                                                                                                                                                                                                                          7.17e+03
38.00
87.50%
87.50%
3.44%
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75.00%
75.00%
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ZIP: 77010-3095
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Percent Similarity:
                                                                                                                                                                                                                                                           Alignment Scores:
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PCT-US95-06987-3
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GENERAL INFORMATION:
APPLICANT: Reed, John
TITLE OF INVENTION:
ANTISENSE OLIGONUCLEOTIDES FOR INHIBITING THE
TITLE OF INVENTION: GROWTH OF CELLS EXPRESSING THE HUMAN BCL-2 GENE
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
ADDRESSEE: P.C.
STREET: 224 Airport Parkway
                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Kiefer, Michael C.
APPLICANT: Kiefer, Michael C.
APPLICANT: OSSINa, Natalya K.
TITLE OF INVENTION: Bak PROMOTER EXPRESSION SYSTEM
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/367,750
FILING DATE: 07-DEC-1999
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative:
Mismatches:
                                                                                                                                                                                                                                                            E: LXR BIOTECHNOLOGY INC.
3095 Richmond Parkway, Suite 213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-375-514-22 (1-205) x US-09-367-750-5 (1-36)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Brown, Theresa A.
REGISTRATION NUMBER: 32,547
REFERENCE/DOCKET NUMBER: 4147-14-PUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           135 GluGluLeuPheArgAspGlyValAsnTrp 144
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                                                                         US-09-367-750-5

Sequence 5, Application US/09367750

Patent No. 6436639
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; Sequence 3, Application US/08217082A
; Patent No. 5734033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 863-9700
TELEFAX: (303) 863-923
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9.17e+03
38.00
70.00%
60.00%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid
EDNESS: single
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STREET: San Jose
                                                                                                                                                                                                                                                                    STREET: 3050
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                              USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                             ZIP: 94806
                                                                                                                                                                                                                                                                ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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GENERAL INFORMATION:

APPLICANT: Reed, John
TITLE OF INVENTION: Regulation of bcl-2 Gene Expression
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
ADDRESSEE: P.C.
STREET: 1755 S. Jefferson Davis Hwy., Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             00000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,485A FILING DATE: 05-JUN-1995
                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                       ATTORNEY, MAINT MANUEL MANUEL MANUEL FORTLORY, AND MANUER 34,600
REGISTRATION NUMBER: 34,600
REGISTRANCE/COCKET NUMBER: 3335-067-55 FWC
TELEPHONE: (408) 436-2070
TELEPHONE: (408) 436-2075
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-375-514-22 (1-205) x US-08-217-082A-3 (1-20)
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Matches:
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                                                                                                                                              APPLICATION NUMBER: US/08/217,082A APPLICATION NUMBER: 24-MAR-1994
CLASSIFICATION: 435
PRICH APPLICATION: 435
PRICH APPLICATION DATA: APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
PRICH APPLICATION DATA: APPLICATION NUMBER: US 07/288,692
FILING DATE: 22-DEC-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: other nucleic acid DESCRIPTION: Synthetic DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BSULT 50
|S-08-465-485A-3/c
Sequence 3, Application US/08465485A
Patent No. 5831066
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100.00%
100.00%
3.35%
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                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Arlington
STATE: Virginia
COUNTRY: U.S.A.
ZIP: 22202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Percent Similarity:
Best Local Similarity:
Query Match:
U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alignment Scores:
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  COUNTRY:
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REGISTRATION NUMBER: 34,600
REFERENCE/DOCKET NUMBER: 3335-070-55 CONT
TELEPHONE: (408) 436-2070
TELEPHONE: (408) 436-2070
INFORMATION FOR SEQ ID NO: 5EQUENCE CHARACTERISTICS: LENGTH: 20 base pairs
TENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-375-514-22 (1-205) x US-08-465-485A-3 (1-20)
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Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/124,256
FILING DATE: 20-SEP-193
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 77/840,716
FILING DATE: 21-FEB-192
PRIOR APPLICATION PUTS:
APPLICATION NUMBER: SPEC-1988
FILING DATE: 22-DEC-1988
ATTORNEY AGENT: INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Search completed: May 31, 2003, 23:23:16 Job time : 67 secs
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MOLECULE TYPE:
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pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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-Q-CGGR2_1/USFTO_spool/US09375514/runat_28052003_165345_18660/app_guery.fasta_1.391
-Q-CGGR2_1/USFTO_spool/US09375514/runat_28052003_165345_18660_LODGEXT=0
-DB-GenEmbl.OFWT=fastap -SUFFIX=rge-MINMATCH=0.1 -LAOPTCH=50
-UNITS=bits -STRAT=1 -END=1 -MATRIX=blosum62 -TRRANS=human40.cdi -LIST=500
-UNITS=bits -STRAT=1 -END=1 -MATRIX=0 -MATISN=0 -ALIGN=50 -MODE=LOCAL
-DOTEWT=pto -NORM=ext -HEAPSIZE=500 -MINLEN 10 -MAXIT -BSBLOCK=100 -LONGLOG
-NO_THEOUT=120 -NORM=EXT -NEGSGOUERY -NEG SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG
-NO_THEOUT=120 -WARN-TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-PGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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                                                                                                                                       May 31, 2003, 18:42:35 ; Search time 1764 Seconds
                                GenCore version 5.1.5
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                        OM protein - nucleic search, using frame_plus_p2n model
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Listing first 500 summaries
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AX031294 Sequence AX031320 Sequence

AR080355 Sequence AR148310 Sequence

Sequence

Sequence

AJ005138 Homo sapi

AR007294 Sequence

Sequence 4

AR020782 Sequence AR052606 Sequence

Sequence Sequence Sequence

AX377550 Sequence

160569 Sequence 21

AX022526 Sequence

AX248864 Sequence AX477731 Sequence AX477732 Sequence

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AX097444 Sequence AX002261 Sequence AX080990 Sequence AX2488293 Sequence AX22498 Sequence AX224989 Sequence

AX262441 Sequence AX248386 Sequence

AR105290 Sequence AX17528 Sequence AX298019 Sequence AX463304 Sequence AX463300 Sequence AX067855 Sequence AX067855 Sequence AR18073 Sequence AR18073 Sequence AR18074 Sequence AX07222 Sequence AX0724243 Sequence AX042445 Sequence AX042445 Sequence AX042445 Sequence AX042445 Sequence AX04245 Sequence AX04245 Sequence AX04245 Sequence AX04245 Sequence AX04249 Sequence AX04245 Sequence AX0436 Sequence AX0436 Sequence AX0436 Sequence AX0368 Sequence AX40381 Sequence AX40381 Sequence AX0368 Sequence AX40381 Sequence AX40381 Sequence AX40381 Sequence AX40381 Sequence AX40381 Sequence AX30787 Sequence AX40381 Sequence AX30787 Sequence AX40381 Sequence AX247608 Sequence AX30787 Seq	
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AX424976 Sequence
I20696 Sequence
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A23776 CASOLII oli
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Muth,J.
Method for detecting mutations in nucleotide sequences
Patent: WO 0212553-A 28 14-FEB-2002;
Nanogen Recognomics GmbH (DE)
Location/Qualifiers
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo
              Sequence
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Mammalia, Eutheria, Primates, Catárrhini, Hominidge, Homo.
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Method for detecting mutations in nucleotide
Patent: WO 0212553-A 27 14-FEB-2002;
Nanogen Recognomics GnbH (DE)
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Mismatches:
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/db_xref="taxon:9606"
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Sequence 28 from Patent W00212553.
AX377551
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Sequence 27 from Patent W00212553.
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VERSION
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PAT 04-DEC-1998
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                                                                                                                                                                                                                                                                                                                                        Unclassified.

1 (bases 1 to 36)
1 (bases 1 to 36)
Thompson,J.D. and Draper,K.G.
Method and reagent for treatment of diseases caused by expression of the bcl-2 gene
Patent: US 5750390-A 1512-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Muth, J.
Method for detecting mutations in nucleotide sequences
Patent: WO 0212553-A 29 14-FEB-2002;
Nanogen Recognomics GmbH (DE)
Location/Qualifiers
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Sequence 15 from patent US 5750390.
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/organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                  Gaps:
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                                                                   1.57e+03
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6.43$
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PAT 29-SEP-1999
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Conservative:
Mismatches:
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Regulation of bcl-2 gene expression
Patent: US 5811066-A 4 03-NOV-1998;
Location/Qualifiers
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Sequence 4 from patent US 5831066.
AR052606.1 GI:5975970
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Sequence 4 from patent US 5734033.
196085.
196085.1 GI:3940555
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AR052606
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                                                                                                                                                                                                                                                                                Unknown.

Inclassified.

(bases 1 to 33)

Thompson,J.D. and Draper,K.G.

Thompson,J.D. and reagent for treatment of diseases caused by expression of the bcl-2 gene
Patent: US 5750390-A 5 12-MAY-1998;
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Guastella,J.
Genes coding for bcl-y a bcl-2 homologue
Patent: US 5789201-A 8 04-AUG-1998;
Location/Qualifiers
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Sequence 8 from patent US 5789201.
AR020782
/organism="Homo sapiens"
./db_xref="taxon:9606"
a 20 c 9 g 3
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Sequence 5 from patent US 5750390.
AR007291 GI:3966775
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/organism="unknown"
16 c 7 g
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9 c 1 g
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AR020782/c
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AR053546/c
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2 (bases 1 to 36)
Kraus, J.P., Le, K., Swaroop, M., Ohura, T., Tahara, T., Rosenberg, L.E., Roper, M.D. and Kozich, V.
Roper, M.D. and Kozich, V.
Splicing and expression in cultured cells.
Human cystathionine beta-synthase CDNA: sequence, alternative Flux. Mol. Genet. 2 (10), 1633-1638 (1993)
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Thompson,J.D. and Draper.K.G.
Method and reagent for treatment of diseases caused by expression
of the bcl-2 gene
Patent: US 5750390-A 8 12-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Gaustadnes,M., Kluijtmans,L.A., Jensen,O.K., Rasmussen,K.,
Heil,S.G., Kraus,J.P., Blom,H.J., Ingerslev,J. and Rudiger,N.
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 Mismatches:
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CBS gene; cystathionine beta-synthase.
Homo sapiens.
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Sequence 8 from patent US 5750390.
AR007294
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/organism="unknown"
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Unclassified.
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Best Local Similarity:
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Best Local Similarity:
Query Match:
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PAT 29-SEP-1999
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Detection of a novel deletion in the cystathionine beta-synthase (CBS) gene using an improved genomic DNA based method FEBS Lett. 431 (2), 175-179 (1998) 98372628
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/organism="Homo sapiens"
/db_xref="taxon:9606"
/chrcmosome="21"
/map="q22.3"
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Sequence 11 from patent US 5834248.
AR053546
AR053546-1 GI:5978408
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O'Reilly, L., Puthalakath, H., Adams, J., O'Connor, L., Cory, S.,
O'Reilly, L., Puthalakath, H., Adams, J., O'Connor, L., Cory, S.,
Novel therapeutic molecules
Patent: WO 9914321-A 16 25-MAR-1999;
INST MEDICAL W & E HALL (AU); PUTHALAKATH HAMSA (AU); REILLY
LORRAINE O (AU); ADAMS JERRY (AU); CONNOR LIAM O (AU)
SUZANNE (AU); HOMG DAVID C S (AU); STRASSER ANDREAS (AU)
LOCATION/QUALIFIERS
                                                                                              1 (bases 1 to 24)
Falb,D.A. and Gimbrone,M.A. Jr.
Compositions and methods for the treatment and diagnosis of
cardiovascular disease using rchd534 as a target
Patent: US 6225084-A 11 01-MAY-2001;
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Sequence 16 from Patent W09914321.
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Sequence 16 from Patent W09914321.
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/organism="unidentified"
/db_xref="taxon:32644"
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AR080355
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Sequence 11 from patent US 5849578.
AR065873.
AR065873.1 GI:5996089
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PAT 20-SEP-2000

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Unclassified.
5 I (bases I to 25)
6 Chenchik A., Jokhadze, G. and Bibilashvilli, R. S. Chenchik A., Jokhadze, G. and Bibilashvilli, R. Methods of assaying differential expression
AL Patent: US 6352829-A 596 05-MAR-2002;
Location/Qualifiers
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Sequence 21 from patent US 5656725.
160569.
160569.1 GI:2479014
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AX022526.1 GI:10046124
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                                                                     Uncases 1 to 28)

O'Reilly, L., Puthalakath, H., Adams, J., O'Connor, L., Cory, S., O'Reilly, D.C. and Strasser. A.

Novel therapeutic molecules
Patent: WO 9914321-A 25-MAR-1999;
INST MEDICAL W & E HALL (AU); PUTHALAKATH HAMSA (AU); REILLY
LORRAINE O (AU); ADAMS JERRY (AU); CONNOR LIAM O (AU); CORY
SUZANNE (AU); HUANG DAVID C S (AU); STRASSER ANDREAS (AU)
Location/Qualifiers
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Uncatasified.
1 (bases 1 to 25)
Chenchik,A., Jokhadze,G. and Bibilashvilli,R.
Methods of assaying differential expression
Patent: US 5994076-A 596 30-NOV-1999;
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/organism="unidentified"
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11 c 4 q
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              AX031320.1 GI:10278645
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PAT 07-SEP-2000
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Unclassified.
1 (bases 1 to 39)
Chittenden,T.D. and Lutz,R.J.
Peptides and compositions which modulate apoptosis
Peptides and compositions which modulate apoptosis
Petent: US 5656725-A 21 12-AUG-1997;
Location/Qualifiers
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PAT 10-AUG-1998

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symma delta T cell antigen receptor delta-chain [V delta 2-J delta 1 junction] [human, skin lesion, Genomic, 33 nt].
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Uyemura,K., Klotz,J., Pirmez,C., Ohmen,J., Wang,X.H., Ho,C., Microanatomic clonality of gamma delta T cells in human leishmaniasis lesions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /gene-"gamma delta T cell antigen receptor delta-chain" 7 c 10 g 6 t
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Location/Qualifiers
                                                                                                      linear
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Yamaguchi, W, Kojima,T., Oh-eda,M. and Hattori,K.
DNA coding for meglakaryocyte potentiator
Patent: US 572318-A 28 03-MAR-1998;
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Mismatches:
Indels:
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Matches:
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189980
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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Admas, J. M., Holmgreen, S. P., Cory, S. and Gibson, L. M.
A novel mammalian gene, bcl-w, belongs to the bcl-2 family of
apoptosis-controlling genes
Patent: WO 9735971-A 3 02-0CT-1997;
ADAMS JERRY MCKEE (AU); HOLMGREEN SHAUN P (AU); CORY SUZANNE (AU);
GIBSON LEONIE M (AU); AMRAD OPERATIONS PTY LTD (AU)
Location/Qualifiers
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          Adams, J.M., Holmgreen, S.P., Cory, S. and Gibson, L.M.
A novel mammalian gene, bcl-w, belongs to the bcl-2 family of apoptosis-controlling genes
Patent: EP 0932674-A 3 04-AUG-1999;
AMRAD OPERATIONS PTY LTD (AU)
Location/Qualifiers
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PAT 20-APR-2002

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Integrated: 1 (Dases 1 to 38)
Rando, R.F., Fennewald, S., Zendegui, J.G., Ojwang, J.O., Hogan, M.E., Pommier, Y. and Mazumder, A.
Guanosine-rich Oligonucleotide integrase inhibitors
Patent: US 6355785-A 112-MAR-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unclassified.

1 (bases 1 to 38)
Rando, R.F., Fenneyald, S., Zendegui, J.G. and Ojwang, J.O. Anti-viral guanosine-rich oligonucleotides
Patent: US 5567604-A 2 22-0CT-1996;
Location/Qualifiers
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Sequence 2 from patent US 5567604.
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Sequence 2 from patent US 6355785.
AR200245 GI:20250319
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                        US-09-375-514-22 (1-205) x AR168776 (1-38)
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Rando,R.F., Ojwaug,J.O., Hogan,M.E., Wallace,T.L. and Cossum,P.A. Anti-viral guanosine-rich tetrad forming oligonucleotides
Patent: US 6288042-A 2 11.5EP-2001;
Location/Qualifiers
                                                                                                                                                                    linear
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Meagher, R.B. and Summers, A.O.
Metal resistance sequences and transgenic plants
Patent: US 5668294-A 6 16-SEP-1997;
Location/Qualifiers
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Sequence 6 from patent US 5668294.
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Sequence 2 from patent US 6288042.
AR168776 GI:17904871
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PAT 06-FEB-1997

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PAT 06-MAY-1999
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Modriquez, R.P., Valladares, J.L. and Mateo, D.A.
Method for obtaining modified immunoglobulins with reduced
immunogenicity of murine antibody variable domains, compositions
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Cornelissen, A.W. and Schallig, H.D.
VACCINE AGAINST GASTRO-INTESTINAL NEWATODES
Patent: WO 9801550-A 16 15-JAN-1998;
UNIV UTRECHT (NL)
Location/Qualifiers
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Matches:
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Patent: EP 0699755-A 10 06-MAR-1996;
INMUNOLOGIA MOLECULAR CENTRO (CU)
Other publication CA.2153135 951231.
Location/Qualifiers
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Sequence 16 from Patent W09801550.
A68670
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Sequence 10 from Patent EP0699755.
A47633
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/organism="unidentified"
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1 (bases 1 to 35)

1 (bases 1 to 35)

8 Acdriguez, R. Perez., Mateo de Acosta del Rio,C.Maria. and Valladares,J.Lombardero.
Walladares,J.Lombardero.
Method for obtaining modified immunoglobulins with reduced immunogenicity of murine antibody variable domains, compositions containing them

Location/Qualifiers

Location/Qualifiers
                                                                                                                                                                               1 (bases 1 to 35)
Godfiquez,R.P., Valladares,J.L. and Mateo,D.A.
Method for obtaining modified immunoglobulins with reduced
immunogenicity of murine antibody variable domains, compositions
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Other publication CA 2153135 951231.
Location/Qualifiers
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Patent: EP 0699755-A 8 06-MAR-1996;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

I (bases 1 to 31)

Cargill, M., Ireland, J.S. and Lander, E.S.

Human single nucleotide polymorphisms

Patent: WO 0166800-A 465 13-SEP-2001;
                                                                                                                                                                                                                                                                        Cornali, E., Nehls, M.C., Schlueter, V., Wattler, F., Wattler, S. and
                                                                                                                                                                                                                                                                                                  Non-human animal model for growth deficiency and information processing or cognitive function defects and use thereof Patent: WO 01/2119-A 16 04-0CT-2001; Ingenium Pharmaceuticals AG (DE)
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Location/Qualifiers
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Mismatches:
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/db_xref="taxon:32630"
/note="Oligonucleotide"
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/db_xref="taxon:9606"
7 c 8 g 7
                            US-09-375-514-22 (1-205) x AR096669 (1-40)
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                                                         53 ProGlyHisThrProHisPro 59
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synthetic construct
artificial sequences.
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AX262441/c
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AX248386
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                                                                                                                                                          PAT 10-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PAT 08-SEP-2000
                                                                                                                                                                                                                                                                     1 (bases 1 to 36)
Rodriguez, R. Perez., Mateo de Acosta del Rio, C. Maria. and
Valladares, J. Lombardero.
Method for obtaining modified immunoglobulins with reduced
immunogenicity of murine antibody variable domains, compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 40)
Garfinkel L. and Richter, T.
Garfinkel L. and Richter, T.
Dethods of using human von Willebrand factor GPIb binding domain polypeptides
Patent: US 6008193-A 7 28-DEC-1999;
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Patent: US 5712120-A 10 27-JAN-1998;
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Matches:
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Sequence 10 from patent US 5712120.
182277 1 GI:3210574
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Sequence 7 from patent US 6008193.
AR096669
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8 c 14 q
                                        US-09-375-514-22 (1-205) x A68670 (1-36)
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PAT 27-FEB-2001
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1 (bases 1 to 31)
Cargill, M., Ireland, J.S. and Lander, E.S.
Human single nucleotide polymorphisms
Patent: WO 0166800-A 372 13-SEP-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US)
                                                                                                                                                                                                                                                                                                                                   Flt-3 ligand-encoding polynucleotide as a polynucleotide-based
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Mismatches:
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Patent: WO 0109303-A 40 08-FEB-2001;
VICAL INCORPORATED (US)
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Sequence 372 from Patent WO0166800.
AX248293
  Matches:
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Sequence 40 from Patent WO0109303.
AX080990.
AX080990.1 GI:13169927
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/organism="Homo sapiens"
/db_xref="taxon:9606"
a 6 c 17 g
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/organism="synthetic cc
/db_xref="taxon:32630"
/note="DNA Primer"
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synthetic construct
artificial sequences.
1 (bases 1 to 40)
Hermanson, G.G.
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60.00%
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39.00
100.00%
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          Percent Similarity:
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Query Match:
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Best Local Similarity:
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ORIGIN
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AX080990/c
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                                                                                                                                                                                                                                                 synthetic construct.
synthetic construct
artificial sequences.
1 (bases 1 to 35)
Kaplan,H., Entwistle,J., Tanha,J., Narang,S. and Dan,M.
Enhanced phage display libraries of human vh fragments and methods
                                                                                                                                                                                PAT 30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 40)
Bauer, S.Christopher., Abrams, M.Allen., Braford-Goldberg, S.Ruth.,
Caparon, M. Helena., Easton, A. Michael., Klein, B.Kure., McKearn, J.P.
Olins, P.O., Paik, K. and Thomas, J.W.
Methods of using multivariant IL-3 hematopoiesis fusion protein
Patent: US 6361977-A 110 26-MAR-2002;
Location/Qualifiers
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/organism="synthetic construct"
/db_xref="taxon:32630"
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Patent: WO 0118058-A 29 15-MAR-2001;
Novopharm Biotech, Inc. (CA)
Location/Qualifiers
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Sequence 110 from patent US 6361977.
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Sequence 29 from Patent WO0118058.
AX097404
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                                                                     US-09-375-514-22 (1-205) x AX248386 (1-31)
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AR202261.1 GI:20256800
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AR202261/c
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PAT 01-MAY-2001
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1 (Dases 1 to 30)
Morten,J.E.
METHODS FOR ANALYZING LTC4 SYNTHASE POLYMORPHISMS AND DIAGNOSTIC
                                                                                                                                                                                                                                                                                                                                                                              synthetic construct.
synthetic construct
artificial sequences.
1 (bases 1 to 24)
Wu,L., Carey,M.F. and Belldegrun,A.S.
Chimeric transcriptional regulatory element and methods for prostate-targeted gene expression
Patent: WO 0127256.A 54 19-APR-2001;
The Regents of the University of California System (US)
                                                                                                                                                                                                                                                                                                                 linear
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/organism="synthetic construct"
/db_xref="taxon:32630"
/note="SYNTHETIC OLIGONUCLEOTIDE"
3 c 10 g 9 t
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Conservative:
Mismatches:
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Mismatches:
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              1. .20
/organism="Homo sapiens"
/db_xref="taxon:9606"
1 c 6 g C
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Sequence 2 from Patent W09910529.
A98608
398608.1 GI:6781653
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 Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Gray,D.M. and Bollon,A.P.
Libraries of optimum subsequence regions of mrna and genomic dni
for control of gene expression
Patent: WO 0161030-A 142 23-AUG-2001;
Cytoclonal Pharmaceutics, Inc. (US); University of Texas at
Dallas, Dept. of Molecular and Cell Biology (US); Lab. of
Experimental Carcinogenesis, National Cancer Institute/NIH (US)
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Libraries of optimum subsequence regions of mrna and genomic dna for control of gene expression
Patent: WO 0161030-A 143 23-A06-2001;
Cytochonal Pharmaceutics, Inc. (US); University of Texas at Dallas, Dept. of Molecular and Cell Biology (US); Lab. of Experimental Carcinogenesis, National Cancer Institute/NIH (US)
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AX224989
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/db_xref="taxon:9606"
11 c 6 q C
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PAT 12-AUG-2002
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1 (bases 1 to 31)
Cargill,M., Ireland,J.S. and Lander,E.S.
Human single nucleotide polymorphisms
Patent: WO 0166800-A 943 13-SEP-2001;
WHITEHRAD INSTITUTE FOR BIOMEDICAL RESEARCH (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fesik, S.W., Petros, A.M., Yoon, H. and Nettesheim, D.G. Mutant bcl.2 proteins and uses thereof Petent: WO 0240530-A 13 23-MAY-2002; ABBOTT LABORATORIES (US)
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Organism="synthetic construct"

/db_xref="taxon:32630"

/note="Primer"

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Sequence 943 from Patent WO0166800.
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Sequence 13 from Patent WO0240530.
AX477731
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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                         Location/Qualifiers
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US-09-375-514-22 (1-205) x S66556 (1-30)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="This sequence comes from Figure 3; 9 base pair deletion; conceptual translation presented here differs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GenBank staff at the National Library of Medicine created this entry [NCBI glubsq 66556] from the original journal article. This sequence comes from Figure 3.

Location/Qualifiers
Patent: WO 9910529-A 2 04-MAR-1999;
MORTEN JOHN EDWARD NORRIS (GB); ZENECA LTD (GB)
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/protein_id-"AAB20361.1"
/db_xref="GI:239219"
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J. Biol. Chem. 266 (33), 22370-22374 (1991)
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/organism="unidentified"
/db_xref="taxon:32644"
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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linear PAT 29-SEP-1999
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Butaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo.

1 (bases 1 to 38)
Cronin,M.T., Frueh,F. and Brennan,T.M.
Methods for optimizing hybridization performance of polynucleotide probes and localizing and detecting sequence variations
Patent: WO 0166804-A 27 13-SEP-2001;
Protogene Laboratories, Inc. (US)
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Patent: US 5831066-A 3 03-NOV-1998;
Location/Qualifiers
                                                                                                                                        38 bp
Sequence 27 from Patent WO0166804.
AX247606.1 GI:15862295
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Sequence 3 from patent US 5831066.
AR052605
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/organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                60 AlaAlaSerArgAspProValAlaArgThr 69
                                                                              2 GCAGCCTCCAGGATTCCAATTGCCAGGACG 31
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/organism="unknown"
10 c 3 g
                                       US-09-375-514-22 (1-205) x A36722 (1-33)
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Tercero,J.C., Garcia,L.C., Ramos,J.A. and Alemany,J.C. Control of PCR mediated detection of micro-organisms Patent: EP 0586112-A 10 09-MAR-1994;
PHARMA GEN S A (ES)
Other publication JP 7087978 950404.
                                                                                                                                                                                                                                       Fesik, S.W., Petros, A.M., Yoon, H. and Nettesheim, D.G. Mutant bcl.2 proteins and uses thereof Patent: WO 0240530-A 14 23-MAY-2002; ABBOTT LABORATORIES (US)
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/db_xref="taxon:32630"
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/db_xref="taxon:11103"
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Sequence 10 from Patent EP0586112.
A36722
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Sequence 14 from Patent WO0240530.
AX477732
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artificial sequences.
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Gaps: US-09-375-514-22 (1-205) x AR052605 (1-20) DB:

Search completed: May 31, 2003, 22:59:21 Job time: 1776 secs

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2147.254 Million cell updates/sec
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1 MAHAGRIGYDNREIVMKYIH.....HTWIQDNGGWVGASGDVSLG
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

/SIDS2/gcgdata/geneseg/genesegn-embl/NA2000.DAT:*/SIDS2/gcgdata/geneseg/genesegn-embl/NA2001A.DAT::/SIDS2/gcgdata/geneseg/genesegn-embl/NA2001B.DAT::

/SIDS2/gcgdata/geneseg/genesegn-embl/NA1998.DAT:* /SIDS2/gcgdata/geneseg/genesegn-embl/NA1999.DAT:*

	Description	BCL-2 mRNA ribozym BCL-2 mRNA ribozym	bcl-y o	proteolyti	Human Bcl-2 PCR pr Bcl-2 related olig	relate		antise	rino rse P	se primer	primer	se primer fo		Bcl-2 muta	"	mutag	domain re	ig Bak	Bcl-2 mut	-w gene	Primer 3'N for Tn2	neu promo	h ol	egrase eotide	r Bclx 2	produc	Human Bcl-2 mutage vWF GPIb binding d	ure von	Probe for von Will Probe for human vo	umigatus cod	sens	Bcl-2 antisense ol Human Bcl-2 mutage	Bcl-2 mutag	bci-2	C/EBP polyn	ademosime -beta anti	C/EBP polyn	uman adenosine /EBP-beta anti	Human C/EBP polynu	/EBP-beta a	Human C/EBP polynu Human adenosine re	PD 1	C/EBP polyn	Human adenosine re
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72444																																														
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C/EBP-Deta antisen Human C/EBP polynu Human adenosine re C/EBP-beta antisen Human C/EBP polynu	aucii robe j beta	2	nucle	C/EBP-beta antisen Human C/EBP polynu	Human adenosine re	Mouse kappa light Human single nucle	Human Bcl-2 protei	Human BCL-2 prote1 BCl2 RNA RT-PCR nr	BCL-2 PCR primer #	Human BCL-2 PCR pr	Diagnostic primer Human single nucle	Mutagenic primer M	Human bak BH2 doma	Beta actin gene pr	Human WT1 truncati	numan ber-z antise PCR primer for hum	Bcl-2 antisense ol	Brevibacterium lac	FOR PILMET USED TO BC1-2 RNA-DNA hybr	First type III-C a	Fibronectin iragme Fibronectin PCR 3'	Human fibronectin	C/EBP-beta antisen Human C/EBP polynu	Human adenosine re	OST311 PCR primer C/EBP-beta antisen		Human adenosine re	C/EBF Deta antisen Human C/EBP polynu	Human adenosine re	Human single nucle Human single nucle	Spider silk protei	Human IL-17RH2 hyb Chimeric BGS prote	Cricetulus griseus	HIV4par, directed	Triplex forming of	Sequence of synthe	Sequence of contro	Triple helix-formi	Triple helix-formi	Guanosine rich oli Viral integrase in	Oligonucleotide #3	BCG deletion regio	Polynucleotide sea	Oligonucleotide 33	ybridis	Synthetic KNA Sequ Antisense oligonuc
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apture oligon LA DOB gene P LA	HTAT Sequence-spec P. furiosus PI-Pfu 5' PCR primer to a Nucleotide fragmen Insecticidal gene Human genomic DNA Human penomic DNA Human ingle nucle Human single single Human membrane-ass Human membrane-ass Human Miller-Dieke Sequence binding t	NGFRSpar, targette Enzymatic RNA mole PCR primer A from Linker #2 used in Llama sdab gene PC Encodes linker pep Permutein linker e PKCalpha primer pa Human PDE7Al 3' UT Hepatitis C virus Human PRO1107 hybr Probe used to dete Human PRO1107 hybr Human PRO1107 hybr Human PRO1107 dete Human NGGO G-Cleav Adipocyte fatty ac C-myc gene target Human beta-globin Beta actin gene pr	uman manganese uman beta-glob uman Inverted uman V gene li -galactono-1,4 BV detecting p NA encoding ME ynthetic TMP g uclear export unan Inverted ligonucleotide ligonucleotide cobe COD 987 s equence of CDR -kit mRNA anti
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Multiple drug resistance; mdr-1; ribozyme; membrane protein; liver; resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin D; vinblastine; small intestine; kidney; adrenal gland; adenocarcinoma; bowel; transformed phenotype; promyelocytic leukemia; human; chronic myelogenous leukemia; CML; follicular lymphocytic leukemia; be-cell acute lymphocytic leukemia; brast cancer; colon carcinoma; neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif; hairpin; hepatitis delta virus; group I intron; RNasep; ss.
                                                                  BCL-2 mRNA ribozyme cleavable nucleotide (1997).
           AAQ51961 standard; RNA; 36 BP
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92US-0882885
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                                               26-MAY-1994 (first entry)
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                                                                                                                                                                         Homo sapiens.
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AAQ51961
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Claim 3; Fig 6; 69pp; English.

New enzymatic RNA molecules (ribozymes) - which cleave mRNA associated with tumours or mRNA expressed from gene encoding

multiple drug resistance

sequences given in AAQ51825-2266 represent areas of mRNAs which are associated with development or maintenance of chronic myelogenous leukemia (CML), promyelocytic leukemia, Burkitt's lymphoma, or acute lymphocytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, breast cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAs containing these target sequences, encode aberant cellular proteins which are able to control cellular proliferation and are directly linked to a leukemic phenotype. These target sequences are identified by the ribozyme of the invention. The ribozymes is formed in a hammerhead motif, but may also be formed in the motif of a hairpin, hepatitis delta virus, group I introm or RNasep-like RNA. These ribozymes and be used to inhibit other animals by modulating expression of the corresponding gene. Cleavage of target mRNAs expressed in pre-neoplastic and transformed cells elicits inhibition of the transformed state. Multiple drug resistance (mdr-1) mRNA specific ribozymes remove the mechanism of drug resistance used by transformed cells and thus enhances drug therapies for tumnours. The ribozymes may also be used to study the development or expression of a transformed phenotype in man and genetic drift and mutations within cells.

Sequence 36 BP; 10 A; 13 C; 8 G; 5 U; 0 other;

Aliqnment Scores:

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Multiple drug resistance; mdr-1; ribozyme; membrane protein; liver; resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin D; vinblastine; small intestine; kidney; adrenal gland; adenocarcinoma; bowel; transformed phenotype; promyelocytic leukemia; human; chronic myelogenous leukemia; CML; follicular lymphocytic leukemia; becall acute lymphocytic leukemia; acute lymphocytic leukemia; brast cancer; colon carcinoma; neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif; hairpin; hepatitis delta virus; group I intron; RNaseP; ss.
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                     Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                              BCL-2 mRNA ribozyme cleavable nucleotide (1592).
            Matches:
                                                                                                181 LeuAsnArgHisLeuHisThrTrpIleGlnAsp 191
                                                                                                           3 CUGAACCGGCACCUGCACACCUGGAUCCAGGAU 35
                                                        Gaps:
                                                                          US-09-375-514-22 (1-205) x AAQ51961 (1-36)
                                                                                                                                                                AAQ51951 standard; RNA; 33 BP.
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92US-0882885.
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                                  Best Local Similarity:
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              Score:
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The sequences given in AAQ51825-2266 represent areas of mRNAs which are associated with development or maintenance of chronic myelogenous leukemia (CML), promyelocytic leukemia, Burkitt's lymphoma, or acute lymphocytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, breast cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAs containing these target sequences, encode aberant cellular proteins which are able to control cellular proliferation and are directly linked to a leukemic phenotype. These target sequences are identified by the ribozyme of the invention. The ribozymes is formed in a hammerhead motif, but may

New enzymatic RNA molecules (ribozymes) - which cleave mRNA associated with tumours or mRNA expressed from gene encoding

(RIBO-) RIBOZYME PHARM INC

Draper KG, Thompson JD; WPI; 1993-386203/48. Claim 3; Fig 6; 69pp; English.

multiple drug resistance

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The primers AAV28335 and AAV28336 were used to amplify the rat bcl-y gene. The mammalian bcl-y genes encode a protein that is a member of the bcl-1 family, components in the cell death pathway. The bcl-2 family have both apoptotic activity and the apoptosis blocking activity. bcl-y falls in the apoptosis activity category. The recombinant protein may administration to recombinant genetic constructs to increase its expression in vivo. Also, antisense constructs can be used in disorders where prevention of cell death is desired.
also be formed in the motif of a hairpin, hepatitis delta virus, group I intron or NaseP-like RNA. These ribozymes may be used to inhibit the development or expression of a transformed phenotype in man and other animals by modulating expression of the corresponding gene.
                                                                Cleavage of target mRNAs expressed in pre-neoplastic and transformed cells elicits inhibition of the transformed state. Multiple drug resistance (mdr-1) mRNA specific ribozymes remove the mechanism of drug resistance used by transformed cells and thus enhances drug therapies for tumours. The ribozymes may also be used to study
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acids encoding B-cell lymphoma-y protein - useful for producing recombinant protein for use in treating uncontrolled cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; bcl-y; bcl-2; cell death pathway; apoptotic; apoptosis; rat; PCR; primer; amplification.
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                                                                                                                                                                                                                                   Length:
Matches:
Conservative:
Mismatches:
                                                                                                                                                                                  Sequence 33 BP; 5 A; 16 C; 7 G; 5 U; 0 other;
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                                                                                                                                   theraples for tumours. The ribozymes may
genetic drift and mutations within cells.
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                                                                                                                                                                                                                                                                                                                                                                                                    3 CCGGGCAUCUUCUCCCCCGGGCCCGGGCAC 32
                                                                                                                                                                                                                                                                                                                                                                                   46 ProGlyIlePheSerSerGlnProGlyHis 55
                                                                                                                                                                                                                                                                                                                                                  (S-09-375-514-22 (1-205) x AAQ51951 (1-33)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV28336 standard; cDNA; 24 BP
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57.00
100.008
100.008
5.168
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970S-0798897
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Best Local Similarity:
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11-FEB-1997;
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Rat bcl-y protein; Rbcl-y; human bcl-y protein; Hbcl-y; bcl-2 homologue; programmed cell death; apoptosis; necrosis; cell death inhibitor; stroke; head trauma; Alzheimer's Disease; neural; muscular degenerative disease; multiple sclerosis; myocardial infarction; vitally induced cell death; aging; spinal cord injury; amyotrophic lateral sclerosis; cancer; premature cell death; cell death; stimulator; prolonged cell life span; Kaposi's sarcoma; lung cancer; autoimmune; hyperimmune disease; parasite; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primers AAX15947-48 were used to amplify cDNA encoding rat bcl-y protein (Rbcl-y). The specification also describes human bcl-y protein (Hbcl-y). Rbcl-y and Hbcl-y are homologues of the bcl-2 protein thought to be involved in programmed cell death (apoptosis and necrosis). Rbcl-y and Hbcl-y proteins may be used to treat conditions associated with a disruption of the cell death pathway. If they act as cell death almohibitors, they may be used in therapies to treat subjects suffering from: strokes, head trauma, Alzheinmer's Disease, neural and muscular degenerative diseases (especially multiple sclerosis), myocardial infarction, vitally induced cell death, aging, spinal cord injuries and amyotrophic lateral sclerosis- conditions where cells under go premature cell death as a result of triggers which may or may not be apparent. They may also be used in this way to develop cell lines which remain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            - useful
                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer used to amplify cDNA encoding the rat bcl-y protein.
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Matches:
Conservative:
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Sequence 24 BP; 3 A; 9 C; 1 G; 5 T; 6 other;
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                                                                                                                                                                                                         188 TrpileGlnAspAsnGlyGlyTrp 195
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97US-0798897.
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100.00%
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4.98%
19
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ID AAX15948 standard; DNA; 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-214150/18.
                                                                                                   Best Local Similarity:
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viable in culture for an extended period. In contrast, if they act as eachl death stimulators, Rbcl-y and Hbcl-y may be used to treat conditions associated with prolonged cell life span such as cancer (especially kaposi's sarcoma and lung cancer) and auto/hyperimmune diseases. They may also be used to cause cell death in, and hence control, parasites.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to human proteolytic enzyme regulatory protein 12 with cytostatic, virucidal, immunomodulatory, antiinflammatory and mamostatic activity. The polypeptide and encoded polynucleotide are applicable in diagnosis and treatment of malignant tumnour, haemopathy, HIV infection, immunological diseases and various inflammations. The present sequence is that of a human proteolytic enzyme regulatory protein 12 PCR primer, useful to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; proteolytic enzyme regulatory protein 12; cytostatic; virucidal; immunomodulatory; antiinflammatory; haemostatic; malignant tumour; HIV; infection; human immunodeficiency virus; immunological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New human proteolytic enzyme regulatory protein 12 and encoded bolynucleotide, applicable in diagnosis and treatment of cancer, haemopathy, human immunodeficiency virus infection, immunological diseases and inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human proteolytic enzyme regulatory protein 12 PCR primer 3.
                                                                                                                                                   480000
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                                                                                                                                                Length:
Matches:
Conservative:
Mismatches:
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                                                                                                  Sequence 24 BP; 3 A; 9 C; 1 G; 5 T; 6 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length:
Matches:
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                                                                                                                                                                                                                                 Gaps:
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                                                                                                                                                                                                                                                             US-09-375-514-22 (1-205) x AAX15948 (1-24)
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                                                                                                                                                                                                                                                                                             188 TrplleGlnAspAsnGlyGlyTrp 195
                                                                                                                                                                                                                                                                                                              5.66e+03
55.00
100.00%
100.00%
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50.00
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                                                                                                                                                                                                                                                                                                                                                                                          AAI64676 standard; DNA; 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-DEC-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; proteolytic enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-597100/67.
                                                                                                                                                                              Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200172793-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mao Y, Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens,
                                                                                                                               Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alignment Scores:
Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-OCT-2001,
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Human; Bcl-2; gene therapy; apoptosis inhibitor; mutant; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to an apoptosis inhibitor comprising the amino acid sequence of Bcl-2 protein in which at least one serine residue is substituted by alanine or aspartic acid. The protein has increased apoptosis inhibitory activity compared with the wild type Bcl-2 protein. The mutated Bcl-2 protein is useful in the treatment of disorders caused by apoptosis. The present sequence was used to amplify DNA encoding the human Bcl-2 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New apoptosis inhibitors, useful for treating apoptosis related
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Mismatches:
Indels:
   Conservative:
Mismatches:
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Matches:
                              Indels:
                                        Gaps:
                                                               US-09-375-514-22 (1-205) x AAI64676 (1-33)
                                                                                      181 LeuAsnArgHisLeuHisThrTrpIle 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; Page 39; 43pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 MetAlaHisAlaGlyArgThrGlyTyr 9
                                                                                                     31 TTAAATAGGCATACACACACAGGATC
                                                                                                                                                                                                                                        Human Bcl-2 PCR primer oligo-10.
                                                                                                                                                                                                                                                                                                                                                                                                                      (HISM ) HISAMITSU PHARM CO LID.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABN85362 standard; DNA; 24 BP
                                                                                                                                                                                                                                                                                                                                                                     2000WO-JP08667.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.85e+04
50.00
100.00%
100.00%
 88.89%
88.89%
4.53%
                                                                                                                                                                                                                (first entry)
                                                                                                                                                               AAH45307 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Shibazaki F, Kuma H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-381681/40.
Percent Similarity:
Best Local Similarity:
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                    07-DEC-2000;
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                                                                                                                                                                                                                  10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alignment Scores:
                                                                                                                                                                                                                                                                                                                                           14-JUN-2001
                                                                                                                                                                                         AAH45307;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disorders
                        Query Match:
DB:
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ABN85362/c
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№
                                                                                                                                          RESULT
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ABN85362;

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20-SEP-1994;
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           RU2181773-C2
                                                                                                                                                                                                                                                                                                                                                                                                                               Alignment Scores:
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                                   27-APR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ86646;
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9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score:
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                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method for estimating transcription levels of genes encoding interferons, interferon-dependent and proliferative cytokines. The method involves determining cytokine mRNA levels, using a combination of RT-PCR with blot- and dot-hybridisation. To illustrate the method, cytokines (alpha-1, beta-1 and gamma-1 interferon), interferon-regulated enzymes (2',5'-oligoadenylatesynthetase, RNABase L and ds-protein kinase) and factors of cellular proliferation (Bcl-2, Ras antigen and gamma-actin cytokeleton protein) were used. The present oligonucleotide was used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Interferon; cytokine; alpha-1 interferon; beta-1 interferon; Bcl-2; gamma-1 interferon; interferon-regulated enzyme; ds-protein kinase; 2',5'-oligoadenylatesynthetase; RNAase L; Fas antigen; gamma-actin cytoskeleton protein; ss.
                                                              Interferon; cytokine; alpha-1 interferon; beta-1 interferon; Bcl-2;
gamma-1 interferon; interferon-regulated enzyme; ds-protein kinase;
2',5'-oligoadenylatesynthetase; RNAase L; Fas antigen;
                                                                                                                                                                                                                                                                                                                                                    Method of assay of human cytokine status at genetic level
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative:
Mismatches:
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Matches:
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                                                                                                          gamma-actin cytoskeleton protein; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bcl-2 related oligonucleotide #2.
                                         Bcl-2 related oligonucleotide #3.
                                                                                                                                                                                                                                                                     (AMVI-) A MED VIROLOGY RES INST.
                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 7; 10pp; Russian
                                                                                                                                                                                                                  16-MAR-2000; 2000RU-0106253.
                                                                                                                                                                                                                                            16-MAR-2000; 2000RU-0106253.
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49.00
100.00%
100.00%
4.44%
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              07-OCT-2002 (first entry)
                                                                                                                                                                                                                                                                                                Sokolova TM, Uryvaev LV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              illustrate the invention.
                                                                                                                                                                                                                                                                                                                        WPI; 2002-391288/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Percent Similarity:
Best Local Similarity:
Query Match:
                                                                                                                                                               RU2181773-C2.
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Pred. No.:
                                                                                                                                     Unidentified.
                                                                                                                                                                                        27-APR-2002.
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interferon), interferon-regulated enzymes (2',5'-oligoadenylatesynthetase, RNAse L and ds-protein kinase) and factors of cellular proliferation (Bcl-2, Fas antigen and gamma-actin cytoskeleton protein) were used. The present oligonucleotide was used to
                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a method for estimating transcription levels of genes encoding interferons, interferon-dependent and proliferative cytokines. The method involves determining cytokine mRNA levels, using a combination of RT-PCR with blot- and dot-hybridisation. To illustrate the method, cytokines (alpha-1, beta-1 and gamma-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Anticode oligomer; antisense oligonucleotide; bcl-2; cancer; therapy; leukemia; lymphoma; solid tumor; breast cancer; autoimmune disease;
                                                                                                                                                                                                                                                        Method of assay of human cytokine status at genetic level
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                227000
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Conservative:
Mismatches:
Indels:
Gaps:
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                                                                                                   (AMVI=) A MED VIROLOGY RES INST.
                                                                                                                                                                                                                                                                                                               Claim 1; Page 7; 10pp; Russian.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     182 AsnArgHisLeuHisThrTrp 188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ86646 standard; DNA; 33 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.68e+04
47.00
100:00%
100.00%
4.26%
16-MAR-2000; 2000RU-0106253.
                                              16-MAR-2000; 2000RU-0106253.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21 AACCGGCACCTGCACACCTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-SEP-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cytoskeleton protein) were illustrate the invention.
                                                                                                                                                       Sokolova TM, Uryvaev LV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bcl-2 splice donor site.
                                                                                                                                                                                                               WPI; 2002-391288/42.
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Best Local Similarity:
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therapeutics
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No.:
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                                                                                                                                                   The antisense oligonucleotide SD-AS (AAQ86645) is complementary to a portion of the splice donor site of the pre-mRNA coding strand of the human bcl-2 gene. It reduces the expression of bcl-2 gene product, thereby inducing programmed cell death of certain cancer cells. The corresp, bcl-2 sense splice donor site region was synthesized for use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     - useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense oligonucleotide; bcl-2 gene; lymphoma; leukaemia; human;
                                          Anti-code oligomers which bind to bcl-2 mRNA - for the treatment of human solid tumours, esp. breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Anti-sense oligo:nucleotide(s) complementary to BCL-2 mRNA for treating cancers, e.g. lymphoma(s) and some leukaemia(s)
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                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                              Sequence 33 BP; 5 A; 5 C; 16 G; 7 T; 0 other;
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-375-514-22 (1-205) x AAQ86646 (1-33)
                                                                                                              Disclosure; Page 13; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human bcl-2 oligonucleotide 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV19654 standard; DNA; 33 BP.
                                                                                                                                                                                                                                                                                                                                                                               2.68e+04
47.00
100.008
100.008
4.268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1998-229881/20.
WPI; 1995-139394/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                              Percent Similarity:
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                                                                                                                                                                                                                                                                      control
                                                                                                                                                                                                                                                                                                                                                             Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV19654;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 10
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Blocking translation at such strategic sites prevents the formation of functional bc1-2 gene product. These oligonucleotides may be used for treating cancers associated with high levels of bc1-2 gene expression, especially lymphomas and some leukaemias.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treating or preventing cancer, tumors and carcinomas, comprises administering B cell lymphoma/leukemia-2 antisense oligonucleotide at high doses for short period for time with one or more cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B cell lymphoma/leukaemia-2; bcl-2; oncogene; antisense; lymphoma; leukaemia; colon carcinoma; rectal carcinoma; pancreatic cancer; breast cancer; ovarian cancer; prostate cancer; renal cell carcinoma; hepatoma; bile duct carcinoma; choriocarcinoma; cervical cancer; testicular cancer; lung carcinoma; bladder carcinoma; melanoma; head and neck cancer; brain cancer; cytostatic; human; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is that of a B cell lymphoma/leukaemia-2 (bcl-2) antisense oligonucleotide. The present invention is directed to the use of bcl-2 antisense oligoners, particularly (33139 (see ABL54148), to treat and prevent bcl-2 related disorders. Administration at high doses results in significant therapeutic responses, including low toxicity, high tolerance and prolonged survival. Administration at high doses for short periods of time (less than 14 days) also provides significant therapeutic responses in the treatment of cancer. The bcl-2 antisense oligomer may also
                                                                                                                                               0000
                                                                                                                                                                                Conservative:
Mismatches:
                                                                                           Sequence 33 BP; 5 A; 5 C; 16 G; 7 T; 0 other;
                                                                                                                                                 Length:
Matches:
                                                                                                                                                                                                                    Indels:
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                                                                                                                                                                                                                                                                                                       Bcl-2 antisense oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                     ABL54152 standard; DNA; 33 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-SEP-2000; 2000US-237009P
10-NOV-2000; 2000US-0709170
                                                                                                                                               .68e+04
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100.00%
100.00%
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                                                                                                                                                                                                                    4.268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GENT-) GENTA INC.
                                                                                                                                                                                                       Best Local Similarity:
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                                                                                                                                                                                  Percent Similarity:
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Multiple drug resistance; mdr-1; ribozyme; membrane protein; liver; resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin D; vinblastine; small intestine; kidney; adrenal gland; adenocarcinoma; bowel; transformed phenotype; promyelocytic leukemia; human; chronic myelogenous leukemia; CML; follicular lymphocytic leukemia; breall acute lymphocytic leukemia; breal acute lymphocytic leukemia; breast cancer; colon carcinoma; neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif; hairpin; hepatitis delta virus; group I intron; RNaseP; ss.
           therapeutics, and in combination with hormone treatment or gene therapeutics, and in combination with hormone treatment or gene cancer of the haematopoietic system, skin, bone and soft tissue, reproductive system, paritourinary system, breast, endocrine system, brain, central nervous system, peripheral nervous system, kidney, lung, respiratory system, thorax, gastrointestinal and alimentary canal, lymph nodes, pancreas, hepatobiliary system, or lymphoma, leukaemia, colon carcinoma, rectal carcinoma, pancreatic, brast, ovarian, prostate, cervical, testicular, head and neck or choract, canal call carcinoma, bile duct carcinoma, choractocarcinoma, lung carcinoma, hepatoma, bile duct carcinoma, choractocarcinoma, lung carcinoma, bladder carcinoma and melanoma
to increase the sensitivity of a subject to cancer
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                                                                                                                                                                                                                                                                                                            Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                               Length:
Matches:
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920S-0936531.
920S-0936532.
920S-0987131.
930S-0006122.
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47.00
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                                                                                                                                                                                                                                                                                                           ercent Similarity:
    est Local Similarity:
    Query Match:
                                                                                                                                                                                                    (all claimed).
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Pred. No.:
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The sequences given in AAQ51825-2266 represent areas of mRNAs which are associated with development or maintenance of chronic myelogenous leukemia (CML), promyelocytic leukemia, Burkitt's lymphona, or acute lymphocytic leukemia, follicular lymphona, B-cell acute lymphocytic leukemia, breast cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAs containing these target sequences, encode aberant cellular proteins which are able to control cellular proliferation and are directly linked to a leukemic phenotype. These target sequences are identified by the ribozyme of the invention. The ribozymes is formed in a hammerhead motif, but may also be formed in the motif of a hairpin, hepatitis delta virus, group in intro or RNasep-like RNA. These ribozymes may be used to inhibit the development or expression of a transformed phenotype in man and other animals by modulating expression of the corresponding gene. Cleavage of target mRNAs expression of the corresponding gene. Cleavage of target mRNAs expressed in pre-neoplastic and transformed cells elicits inhibition of the transformed state. Multiple drug resistance (mdr-1) mRNA specific ribozymes remove the mechanism of therapies for tumours. The ribozymes remove the mechanism of the rapies for tumours. The ribozymes and also be used to study genetic drift and mutations within cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cardiovascular disease; differential expression; target gene; pathway gene; fingerprint gene; atherosclerosis; ischaemia; reperfusion; hypertension; restenosis; arterial inflammation; vector; antibody; diagnosis; gene therapy; drug screening; bcl-2; polymerase chain reaction; PCR; primer; ss.
                                    New enzymatic RNA molecules (ribozymes) - which cleave mRNA associated with tumours or mRNA expressed from gene encoding
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Matches:
Conservative:
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Indels:
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                                                                                                            Claim 3; Fig 6; 69pp; English.
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ID AAT36054 standard; DNA; 24 BP.
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95US-0386844.
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100.00%
100.00%
4.08%
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                                                                         multiple drug resistance
WPI; 1993-386203/48
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10-FEB-1995;
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(MILL-) MILLENNIUM PHARM INC.

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Fingerprinting gene; rchd502; transmembrane protein; cardiovascular; fingerprint/target gene; up.regulated; endothelial cell; shear-stress; atherosclerosis; ischemia; reperfusion; hypertension; restenosis; human; RT-PCR; primer; bcl-2; mouse; ss.
                                                                                                                                                       Monocyte RNA from apolipoprotein E (apoE)-deficient and control mice was compared using primers for mouse bc1-2 (AAT36053 and AAT36054) and gamma-actin (AAT36055 and AAT36056). bc1-2 mRNA levels were significantly lower in the apoE-deficient mice (an animal model of atherosclerosis). Regulation of the mouse bc1-2 gene appeared to be regulated by serum cholesterol levels. Similar results were obtd. with the human bc1-2 and glutathione peroxidase genes (see also AAT36057-58). The discovery that certain genes (see also AAT36029-36) are differentially expressed in cardiovascular diseases can be used to develop methods for the diagnosis and treatment of such diseases.
                                                                      New genes differentially expressed in cardiovascular disease - and related vectors, host cells, proteins and antibodies, for diagnosis, monitoring, treatment and drug screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Reverse primer for RT-PCR analysis of mouse bcl-2 mRNA.
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Matches:
Conservative:
Mismatches:
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                                                                                                                                 Example 7; Page 12; 200pp; English
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44.00
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95US-0485573
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AAX26250 standard; DNA; 24
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                                           WPI; 1996-384391/38.
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Best Local Similarity:
Query Match:
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              DA;
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02
              Falb
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The invention relates to a rchd502 target/fingerprint gene encoding a transmembrane protein. The invention provides cDNAs contained in plasmids prCHD5025F (ATCC 69982) that encode the rchd502 polypeptide, and are differentially expressed in cardiovascular disease states. Cultured genetically engineered host cell containing the rchd502 polypeptides in operative association with a nucleotide regulatory element are used for producing a polypeptide rchd502 gene product. Identifying that the fingerprint/rarget gene rchd502 is differentially expressed (up-regulated) by endothelial cells subjected to shear stress, provides a tool for the diagnosis and treatment of cardiovascular disease e.g. atherosclerosis, ischemia/reperfusion, hypertension, restenosis. The fingerprint gene is useful for testing the efficacy of candidate drugs in basic research and in clinical trials and or imaging of a diseased cardiovascular tissue. The gene may also be used in screening for ligand-receptor interaction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; cardiovascular disease; atherosclerosis; ischaemia; restenosis; reperfusion; hypertension; arterial inflammation; diagnosis; rchd528;
New polynucleotides consisting of residues 1-1929 of the rchd502 gene - are differentially expressed in cardiovascular disease states, and can therefore be used to treat and diagnose cardiovascular diseases
                                                                                                                                                                                                                                                                                                                                                                                                                               4,0000
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Mismatches:
Indels:
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Matches:
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                                                                                Disclosure; Column 9; 121pp; English.
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95US-0458873
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Query Match:
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07-JUN-1995;
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(MILL-) MILLENNIUM PHARM INC

WPI; 1999-069743/06

Falb DA;

96US-0616844

15-MAR-1996;

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The present invention describes rchd528 protein. A method has been developed for producing the rchd528 gene product. The present invention also describes methods and compositions for the treatment and diagnosis of cardiovascular diseases, including: atherosclerosis; ischaemia; restenosis; reperfusion; hypertension; and arterial inflammation. The present sequence represents a primer used in an example from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel isolated rchd502 polypeptides, differentially expressed in response to endothelial cell shear stress, used for diagnosis, monitoring clinical trails, and treating cardiovascular diseases such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This oligonucleotide is a reverse PCR primer for the mouse bcl-2 gene. It was used with the forward primer given in AAA88586 in a quantitative RT-PCR analysis of mouse bcl-2 mRNA levels in app.8-deficient mice, a murine model of atheroscierosis. bcl-2
           DNA encoding rchd528 polypeptide - associated with cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse; bcl-2 gene; differential expression; atherosclerosis; cardiovascular disease; diagnosis; therapy; PCR primer; ss.
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                                                                                                                                                                                                                                         Length:
Matches:
Conservative:
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                                                                                                                                                                                              Sequence 24 BP; 4 A; 14 C; 1 G; 5 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                        (1-24)
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                                                    Example; Column 95; 122pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BGHM ) BRIGHAM & WOMENS HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer for mouse bcl-2 gene.
                                                                                                                                                                                                                                                                                                                                       US-09-375-514-22 (1-205) x AAV81830
                                                                                                                                                                                                                                                                                                                                                                 140 AspGlyValAsnTrpGlyArg 146
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95US-0386844.
95US-0485573.
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44.00
100.00%
100.00%
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                                                                                                                                                                                                                                                                                                                                                                                 23 GATGGGGTGAACTGGGGGGAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                      24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                       1587/c
AAA88587 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gimbrone MA, Falb DA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-611017/58
                                                                                                                                                                                                                                                                                est Local Similarity:
uery Match:
                                                                                                                                                                                                                                                                   Percent Similarity:
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10-FEB-1995;
07-JUN-1995;
                                                                                                                                                                                                                          Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US6124433-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ischemia
                                                                                                                                                                    invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAA88587;
                           disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus sp.
                                                                                                                                                                                                                                          Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA88587,
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This sequence represents a PCR primer used to amplify the mouse bcl2 nucleotide sequence. The primer is used in methods for the flachtification of the marker proteins and differentially expressed genes of the invention. The invention relates to the rchd523 transmembrane polypeptide (see AAY78506) encoded by cDNA contained in the plasmid pfchd523. The rchd523 protein is differentially expressed in diseased cells compared to healthy cells. The rchd523 protein may be used as a marker protein for the diagnosis of cardiovascular diseases including atherosclerosis, ischaemia, reperfusion, hypertension, restenosis and arterial inflammation. Trchd523 peptides may be used as antigens in the production of antibodies specific for rchd523. The anti-rchd523 antibodies may then be used in diagnostic assays to quantitate rchd523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Differentially expressed; cardiovascular disease; atherosclerosis; ischaemia; reperfusion; hypertension; restenosis; arterial inflammation; bcl2; differential display analysis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Marker proteins for the diagnosis of cardiovascular diseases such as atherosclerosis and hypertension, comprising peptide sequences derived from the rchd523 transmembrane protein -
mRNAs were shown to be lower in apoE-deficient mice relative to wild type controls. The invention provides novel human genes (see AAAA86576-83) that are differentially expressed in cardiovascular disease states, and which are of diagnostic and therapeutic use.
                                                                                                                                    4,0000
                                                                                                                                                                   Conservative:
Mismatches:
                                                                               4 A; 14 C; 1 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Reverse primer for mouse bcl2 amplification.
                                                                                                                                    Length:
Matches:
                                                                                                                                                                                                      Indels:
                                                                                                                                                                                                                                                      (1-24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Column 9; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BGHM ) BRIGHAM & WOMENS HOSPITAL. (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                      US-09-375-514-22 (1-205) x AAA88587
                                                                                                                                                                                                                                                                                      140 AspGlyValAsnTrpGlyArg 146
                                                                                                                                                                                                                                                                                                         23 GATGGGGTGAACTGGGGGGGG 3
                                                                                                                           3.06e+04
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95US-0485573.
                                                                                                                                                                                                                                                                                                                                                                                       AAZ89800 standard; cDNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        97US-0944423.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Falb DA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-146911/13.
                                                                                                                                                                   Percent Similarity:
Best Local Similarity:
                                                                                 Sequence 24 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-0CT-1997;
                                                                                                                    Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6020463-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ89800;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mus sp.
                                                                                                                                                                                                      Query Match:
                                                                                                                                                                                                                                                                                                                                                                    AAZ89800/c
                                                                                                                                                                                                                                                                                                                                                      RESULT 17
 8888888
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peptides in samples.

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The present invention describes an isolated polypeptide (I) comprising either the amino acid sequence of 1481 residues, given in AAK6847, or an amino acid sequence encoded by the cDNA contained in plasmids prCHD528A (ATCC 69985), prCHD528B (ATCC 69986) and prCHD528C (ATCC 69987). The polypeptide is useful in the treatment and diagnosis of cardiovascular disease, such as, atherosclerosis, ischaemia/reperfusion, hypertension, restenosis and arterial inflammation. AAZ88001 to AAS88040, and AAX68444 to AAX68457 represent sequences used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated polypeptide for treating and diagnosing cardiovascular disease, such as, atherosclerosis, ischemia/reperfusion, hypertension, restenosis and arterial inflammation -
                                                                                                                                                                                                                                                                                                                                                Cardiovascular disease; diagnosis; atherosclerosis; ischaemia; reperfusion; hypertension; restenosis; arterial inflammation; antiarteriosclerotic; vasotropic; hypotensive; PCR primer; ss.
                                        40000
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                                                                Conservative:
Mismatches:
Indels:
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5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 24 BP; 4 A; 14 C; 1 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                        Mouse bcl-2 reverse PCR primer SEQ ID NO:11.
                                    Length:
Matches:
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                                                                                                         Gaps:
                                                                                                                                  (1-24)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MILL-) MILLENIUM PHARM INC. (BGHM ) BRIGHAM & WOMENS HOSPITAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example; Column 9; 122pp; English.
A; 14 C; 1 G;
                                                                                                                               US-09-375-514-22 (1-205) x AAZ89800
                                                                                                                                                        AAZ88011 standard; DNA; 24 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96US-0599654.
95US-0386844.
95US-0485573.
                                    3.06e+04
44.00
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                                                                                                                                                                                                                                                                                            19-APR-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Falb DA, Gimbrone MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-136704/12.
                                                                          Best Local Similarity:
Query Match:
DB:
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Sequence 24 BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Percent Similarity:
                                                                Percent Similarity:
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10-FEB-1995;
07-JUN-1995;
                          Alignment Scores:
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                                                                                                                                                                                                                                                                   AAZ88011;
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                                                     Score:
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The present invention describes methods for inhibiting the growth of (1), inducing apoptosis in (II), inhibiting tumourigenesis in (III), inhibiting neoplastic transformation in (IV) cancer cells and for enhancing the effectiveness of chemo-(IV) and radiotherapies (VI) for the treatment of cancer. The methods comprise administering an antagonist of STAT (signal transducer and activator of transcription) signalling. The methods may be used for inhibiting the growth of cancer cells (II), inhibiting tumourigenesis in cancer cells (II), inhibiting tumourigenesis in cancer cells (III), inhibiting neoplastic transformation in cancer cells (IV) and for enhancing the effectiveness of chemo-(IV) and radiotherapies (VI) for the treatment of cancer. The present sequence represents a specific gene amplification PCR primer for boll-2, which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Administering antagonists of STAT (signal transducer and activator of transcription) signaling in cells for the treatment of cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STAT; signal transducer and activator of transcription; human; cell signalling; cytokine; growth factor; oncogenesis; tumour; apoptosis; cytostatic; tumourigenesis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Heller R, Jaroszeski M;
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Matches:
Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bcl-2 specific gene amplification PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 25 BP; 3 A; 13 C; 5 G; 4 T; 0 other;
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   Indels:
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                               Gaps:
                                                                                              US-09-375-514-22 (1-205) x AAZ88011 (1-24)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 9; Page 31; 92pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       102 AspAspPheSerArgArgTyrArg 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 GACGACTICICCCGCCGCTACCGC 25
                                                                                                                                                                 146
                                                                                                                                                                                                                                                                                                                                                                     BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JAN-2000; 2000WO-US01845.
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44.00
100.008
100.008
3.998
                                                                                                                                                                 140 AspGlyValAsnTrpGlyArg
                                                                                                                                                                                                                                23 GATGGGGTGAACTGGGGGAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-DEC-2000 (first entry)
3.99%
21
                                                                                                                                                                                                                                                                                                                                                                 AAA73296 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-505964/45.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200044774-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA73296;
   Query Match:
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                                                                                                                                                                                                                                                                                                  RESULT 19
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Chenchik A,
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No.:
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à
                                                                                                                                                                                                                                                                                                                                                                                                                 Human; bcl-2; cancer detection; disseminated cancer cell; cytostatic;
                                                                                      Human; Bcl-2; gene therapy; apoptosis inhibitor; mutant; primer; ss.
                                                                                                                                                                                                                                                                                The invention relates to an apoptosis inhibitor comprising the amino acid sequence of Bc1-2 protein in which at least one serine residue is substituted by alanine or aspartic acid. The protein has increased apoptosis inhibitory activity compared with the wild type Bc1-2 protein. The mutated Bc1-2 protein is useful in the treatment of disorders caused by apoptosis. The present sequence was used to create a mutant Bc1-2 protein of the invention.
                                                                                                                                                                                                                                          New apoptosis inhibitors, useful for treating apoptosis related
                                                                       Human Bcl-2 mutagenic primer oligo-8 for S161A substitution.
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                                                                                                                                                                                                                                                                                                                                                                                        Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                 Sequence 27 BP; 5 A; 12 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                         Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-375-514-22 (1-205) x AAH45305 (1-27)
                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 11; 43pp; Japanese.
                                                                                                                                                                                           (HISM ) HISAMITSU PHARM CO LID.
                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
                                                                                                                                                                                                                                                                                                                                                                    3.46e+04
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100.008
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                      AAH45305 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAL46663 standard; DNA; 27
                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human bcl-2 mRNA probe.
                                                                                                                                                                                                          Shibazaki F, Kuma H;
                                                                                                                                                                                                                          WPI; 2001-381681/40.
                                                                                                                                                                                                                                                                                                                                                                                       Percent Similarity:
Best Local Similarity:
                                                                                                                            WO200142459-A1.
                                                                                                       sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                       10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                Alignment Scores:
Pred. No.:
                                                                                                                                            14-JUN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-AUG-2002
                                                                                                              Synthetic
                                                                                                                                                                                                                                                    disorders
                                       AAH45305;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAL46663;
                                                                                                                                                                                                                                                                                                                                                                                                       uery Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     probe;
     RESULT 20
AAH45305/c
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The present invention relates to a method for the clinical validation of a target from disseminated cancer cells, characterised in that for a population of individuals it is determined whether a status of the target determined in disseminated cancer cells of the individuals correlates with at least one cancer-related information about the clinical status of the individuals. The method is useful for clinically validating target from disseminated cancer cells. The present sequence is a probe used
                                                                                                                                                                                                                                                                                                                                                      Clinically validating target from disseminated cancer cells by determining whether status of target determined in cancer cells of individuals correlates with cancer-related information about clinical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Primer; ss; DNA microarray; differential expression analysis; human.
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Matches:
Conservative:
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Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     demonstrate the method of the invention.
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                                                                                                                                                                                                                                                                 Boeckmann B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; Page 55; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   172 AsnileAlaLeuTrpMetThrGlu 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25 AACATCGCCCTGTGGATGACTGAG
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44.00
100.00%
100.00%
3.99%
                                                                                                 05-NOV-2001; 2001WO-EP12786
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ID ABK66508 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                             status of individuals
                                                                                                                                                                                                                                                                 Siesing M, Grill H,
                                                                                                                                                                                                                                                                                                             WPI; 2002-426739/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity:
                                                                                                                                                                                                                     (GIES/) GIESING
WO200237113-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Percent Similarity:
                                                                                                                                                                      03-NOV-2000;
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                                                                                                                                                03-NOV-2000;
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                                                10-MAY-2002
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WPI; 2002-314699/35

Producing sub-population of labeled nucleic acids, useful for analysing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers

Example 3; SEQ ID No 596; 11pp; English.

The invention relates to producting a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiological source, with a pool of 50 distinct gene specific primers under suitable conditions to enzymatically generate sub-population of NAS, where conditions to enzymatically generated using a single gene specific primer has a sequence complementary to a distinct mRNA, and each labeled NA is generated using a single gene specific primer. The method is useful for producing a sub-population of labeled NAS which is useful for analysing the differences in the RNA profiles comprises producing subpopulation of labeled NAS for the different physiological sources, where the method is preferred by preferred by hybridising the labeled NAS for each of the source to identify differences in the population, where the comparison is preferably performed by hybridising the labeled NAS for each of the sources to an array of probe NAS stably conditions. The sources of a substrate to produce a hybridisation. The sources where differential expression assays are the sources, where differential expression assays are the sources, where differential gene expression assays are tissue e.g. neoplastic a normal tissue, or different tissue or subtissue types. The present sequence is a human gene specific PCR primer used in the method of the invention.

Note: The sequence data for this patent did not form part cof the printed specification, but was obtained in electronic cof the printed specification, but was obtained in electronic cof the printed specification, but was obtained in electronic cof the printed specification, but was obtained in electronic cof the printed specification. http.wipo.seqdata.uspto.gov/sequence.html?DocID=6352829B1.

Sequence 25 BP; 5 A; 11 C; 4 G; 5 T; 0 other

Alignment Scores:

Best Local Similarity: Percent Similarity: Query Match:

Length:
Matches:
Conservative:
Mismatches:
Indels: 4.34e+04 42.00 87.50% 87.50% 3.80%

25 0 0 0 0

US-09-375-514-22 (1-205) x ABK66508 (1-25)

189 IleGlnAspAsnGlyGlyTrpVal 196 ð g

24 ATCCAAGACCAGGGTGGTTGGGTG 1 AAH45298/c RESULT

AAH45298;

AAH45298 standard; DNA; 27

BP

10-SEP-2001 (first entry)

Human Bcl-2 mutagenic primer oligo-1 for S24A substitution.

Human; Bcl-2; gene therapy; apoptosis inhibitor; mutant; primer; ss.

Homo sapiens,

Synthetic.

WO200142459-A1

14-JUN-2001

07-DEC-2000; 2000WO-JP08667

99JP-0350427. 09-DEC-1999;

(HISM) HISAMITSU PHARM CO LTD.

Shibazaki F, Kuma H;

WPI; 2001-381681/40.

New apoptosis inhibitors, useful for treating apoptosis related disorders

Example 1; Page 10; 43pp; Japanese.

The invention relates to an apoptosis inhibitor comprising the amino acid sequence of Bcl-2 protein in which at least one serine residue is substituted by alanine or aspartic acid. The protein has increased apoptosis inhibitory activity compared with the wild type Bcl-2 protein. The mutated Bcl-2 protein is useful in the treatment of disorders caused by apoptosis. The present sequence was used to create a mutant Bcl-2 protein of the invention.

Sequence 27 BP; 5 A; 7 C; 8 G; 7 T; 0 other;

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27
7
1
1
0
0
                Matches:
Conservative:
Mismatches:
        Length:
                                         Indels:
                                                Gaps:
        4.67e+04
                42.00
88.89%
77.78%
3.80%
                        Percent Similarity:
Best Local Similarity:
Alignment Scores:
                                         Query Match:
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JS-09-375-514-22 (1-205) x AAH45298 (1-27)

à 셤

AAH45302/c RESULT 24

BP. AAH45302 standard; DNA; 27

AAH45302;

10-SEP-2001 (first entry)

Human Bcl-2 mutagenic primer oligo-5 for S116A substitution.

Human; Bcl-2; gene therapy; apoptosis inhibitor; mutant; primer; ss.

Homo sapiens. Synthetic.

WO200142459-A1.

14-JUN-2001

07-DEC-2000; 2000WO-JP08667

99JP-0350427 09-DEC-1999;

(HISM) HISAMITSU PHARM CO LTD.

Shibazaki F, Kuma H;

WPI; 2001-381681/40.

New apoptosis inhibitors, useful for treating apoptosis related disorders

Example 1; Page 11; 43pp; Japanese.

The invention relates to an apoptosis inhibitor comprising the amino acid sequence of Bcl-2 protein in which at least one serine residue is substituted by alanine or aspartic acid. The protein has

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increased apoptosis inhibitory activity compared with the wild type Bcl-2 protein. The mutated Bcl-2 protein is useful in the treatment of disorders caused by apoptosis. The present sequence was used to create a mutant Bcl-2 protein of the invention.
       88888
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Sequence 27 BP; 4 A; 8 C; 10 G; 5 T; 0 other;

Length:
Matches:
Conservative:
Mismatches:
Indels: Gaps: 4.67e+04 42.00 100.00% 88.89% 3.80% Percent Similarity: Best Local Similarity: Alignment Scores: Query Match: .. SO:: Pred. Score:

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US-09-375-514-22 (1-205) x AAH45302 (1-27)

112 PheAlaGluMetSerSerGlnLeuHis 120 27 TTCGCCGAGATGGCCAGCCAGCTGCAC 1 ð 셤

ESULT 25

BP. AAT42430 standard; DNA; 39 AAT42430

AAT42430;

29-JUL-1997 (first entry)

GD domain region for Bak encoding DNA, amino acid residues 82-94.

Apoptosis; follicular lymphoma; tumour; p53; antibody; ss.

Synthetic.

WO9635951-A1.

14-NOV-1996.

96WO-US06122. 06-MAY-1996;

95US-0440391. 12-MAY-1995;

(IMMU-) IMMUNOGEN INC.

Chittenden TD, Lutz RJ;

WPI; 1996-518805/51. P-PSDB; AAW06293 Peptide(s) comprising GD domains - have similar activities to wild type Bak, and cause cellular apoptosis for treatment of viral infection

Claim 6; Page 45; 69pp; English.

The term GD domain refers to a protein domain first identified in and shown to be essential for the interaction of Bak with Bcl-x(L) and for Bak's call killing function; and to peptides and/or molecules capable of mimicking its structure and/or function. The present sequence encodes a GD domain corresponding to amino acid residues 82-94 of Bak.

An antibody raised against a GD domain may be used to screen a cDNA cepression library for clones comprising cDNA inserts encoding immunocrossreactive proteins. Truncated GD domain peptides have been shown to maintain the protein binding and cell killing function cannour cell. These peptides act independently of p53 status. Bak or GD domain mimetics that inhibit Bcl.2 may be selectively toxic to certain tumours, e.g. follicular lymphoma, which depend on high levels of Bcl.2 for their continued growth and survival. GD domain mimetics

may also be used for combatting viral infections by causing apoptosis

```
/product= "Bak GD domain region #1"
/note= "This sequence lacks both start and stop codons"
                                                                                                                                                                                            GD domain; apoptosis; interaction with Bcl-XL; cell killing function; bak; cell death regulatory molecule; autoimmune disease; cancer; ds.
                                                                                                                                                                               DNA encoding Bak GD domain region #1 for modulating apoptosis.
                      33
                                     Conservative:
Mismatches:
Sequence 39 BP; 11 A; 11 C; 11 G; 6 T; 0 other;
                                                                                        1 GGGGACGACATCAACCGACGCTATGACTCAGAGTTC 36
                      Length:
Matches:
                                                   Indels:
                                                            Gaps:
                                                                         US-09-375-514-22 (1-205) x AAT42430 (1-39)
                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                   BP
                      6.81e+04
42.00
75.00%
58.33%
                                                                                                                                   ABK11179 standard; DNA; 39
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                                                                                                                                                                                                                                           1..39
/*tag=
                                            Best Local Similarity:
Query Match:
                                     Percent Similarity:
                                                                                                                                                                                                                                                                                       US6221615-B1.
                                                                                                                                                                                                                     Unidentified
                Alignment Scores:
                                                                                                                                                                 05-JUN-2002
                                                                                                                                                  ABK11179;
                                                                                                                     RESULT 26
                                                                                                                            ABK11179
                                Score:
                                                                                         δλ
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Identifying agents (e.g. modulators of apoptosis) capable of modulating GD domain mediated heterodimerisation or homodimerisation comprises carrying out a heterodimerisation or homodimerisation assay

(APOP-) APOPTOSIS TECHNOLOGY INC.

WPI; 2002-234950/29.

Chittenden TD,

P-PSDB; AAU77876.

95US-0440391. 97US-0908597.

12-MAY-1995; 08-AUG-1997; 25-JAN-1999;

99US-0236385.

24-APR-2001.

Disclosure; Fig 8B; 37pp; English.

which are capable of modulating apoptosis. The GD domains are essential for Bak's interaction with Bcl-XL, and to Bak's cell killing function. The GD domains mediate key protein/protein interactions with multiple cell death regulatory molecules. Also described are methods of identifying agonists or antagonists of GD domains. The methods are useful for identifying agents capable of modulating GD domain mediated heterodimerisation or homodimerisation. The methods are particularly useful in drug screening and design, e.g. for identifying agents for treating autoimmune disease or cancer, or for identifying modulators of apoptosis. The present DNA sequence encodes a GD domain region. present invention relates to novel peptides, designated GD domains, The

Sequence 39 BP; 11 A; 11 C; 11 G; 6 T; 0 other;

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AAH45306 standard; DNA; 27
                                                                                                                                                                                                                    Kuma H;
                                                                                                                                                                                                                                   WPI; 2001-381681/40.
                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                       Sequence 27 BP;
                                                                                                                          WO200142459-A1
                                                                                                                                                                                                                                                                                                                                                                                                                  Percent Similarity:
                                                                                                                                                                               09-DEC-1999;
                                                                                                 Homo sapiens
                                                                                                                                                                                                                  Shibazaki F,
                                                                                                                                                                                                                                                                                                                                                                                        Alignment Scores:
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                                           10-SEP-2001
                                                                                                                                            14-JUN-2001
                                                                                                          Synthetic.
                                                                                                                                                                                                                                                               disorders
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                          AAH45306;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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AAH45306,
                                                                                                                                                                                                                                                                                                                                                                                                           Score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene therapy; apoptosis inhibitor; mutant; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an apoptosis inhibitor comprising the amino acid sequence of Bcl-2 protein in which at least one scrine residue is substituted by alamine or aspartic acid. The protein has increased apoptosis inhibitory activity compared with the wild type Bcl-2 protein. The mutated Bcl-2 protein is useful in the treatment of disorders caused by apoptosis. The present sequence was used to create a mutant Bcl-2 protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                       apoptosis inhibitors, useful for treating apoptosis related
                                                                                                                                                                                                Human Bcl-2 mutagenic primer oligo-2 for S24D substitution.
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      Length:
Matches:
Conservative:
Mismatches:
Indels:
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Mismatches:
                                                                                      101 GlyAspAspPheSerArgArgTyrArgGlyAspPhe 112
                                                                                                Sequence 27 BP; 5 A; 6 C; 8 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length:
Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 10; 43pp; Japanese.
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                                                                    US-09-375-514-22 (1-205) x ABK11179
                                                                                                                                            BP.
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41.00
77.78%
77.78%
3.71%
      6.81e+04
42.00
75.00%
58.33%
3.80%
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                                                                                                                                          AAH45299 standard; DNA; 27
                                                                                                                                                                             10-SEP-2001 (first entry)
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                        Percent Similarity:
Best Local Similarity:
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                  Human; Bcl-2;
                                                                                                                                                                                                                                   Homo sapiens.
Synthetic.
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Alignment Scores:
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                                                                                                                                                            AAH45299;
                                          Query Match:
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SS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bcl-w; apoptosis; bcl-2; cell survival; treatment; therapy; cancer; diagnosis; degenerative disease; PCR primer; ss.
                                                                                                      Human; Bcl-2; gene therapy; apoptosis inhibitor; mutant; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New apoptosis inhibitors, useful for treating apoptosis related
                                                 Human Bcl-2 mutagenic primer oligo-9 for S161D substitution.
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Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6 A; 12 C; 4 G; 5 T; 0 other;
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Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 11; 43pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27 ATGTGTGGAGGTCGTCAACCGGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HISM ) HISAMITSU PHARM CO LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
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41.00
88.89%
88.89%
3.71%
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(first entry)
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AAT96580 and AAT96579 are PCR primers used to amplify a novel gene, bcl-w, a member of the mouse bcl-2 gene family. This gene promotes cell survival, so its modulation is useful in treatment of cancer or auto-immune diseases, degenerative diseases (e.g. stroke, Alzheimer's disease, myocardial infarct, muscular degeneration, hypoxia, ischaemia, Upregulation of the gene can also be used to modify cell lines cultured in vivo, e.g. to develop new lines, to facilitate isolation of thybridomas and to increase survival of primary explants during genetic modification. It can be used to produce recombinant Bcl-w for therapy, diagnosis, antibody production or screening of potential modulators.
                                                                                                                                                                                                                                                                                                                                                                                            to induce
                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acid encoding apoptosis related gene bcl-w - used or inhibit cell survival, e.g. for treatment of cancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mercuric ion reductase merA PCR mutagenesis primer 3'A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0000
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Matches:
Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 31 BP; 5 A; 10 C; 5 G; 7 T; 4 other;
                                                                                                                                                                                                                                                                                                                                       Cory S, Gibson LM, Holmgreen SP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-375-514-22 (1-205) x AAT96580 (1-31)
              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 30; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   188 TrpileGlnAspAsnGlyGlyTrp 195
                                                                                                                                                                                                                                                                                                            (AMRA-) AMRAD OPERATIONS PTY LTD.
                                                               'note= "inosine"
                                                                                                                      'note= "inosine"
                                                                                                                                                           /mod_base= i
/note= "inosine"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6.27e+04
41.00
75.00%
3.71%
                                                    /mod base= i
                                                                                                        /mod_base= i
                                                                                                                                                                                                                                                                                 96AU-0008965.
                                                                                                                                                                                                                                                        97WO-AU00199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT92245/c
ID AAT92245 standard; DNA; 36
                                                                                          /*tag- b
                                                                                                                                                 /*tag= c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                        *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                      degenerative diseases
                                                                                                                                                                                                                                                                                                                                                                WPI; 1997-489635/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Percent Similarity:
Best Local Similarity:
                         modified_base
                                                                             modified base
                                                                                                                                 modified_base
                                                                                                                                                                                                   WO9735971-A1
                                                                                                                                                                                                                                                      27-MAR-1997;
                                                                                                                                                                                                                                                                                 27-MAR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lignment Scores:
red. No.:
Score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JAN-1998
                                                                                                                                                                                                                              02-0CT-1997
                                                                                                                                                                                                                                                                                                                                       Adams JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT92245;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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transposon; Tn21; merB; mercuric ion reductase; organomercury lyase; ss; transgenic plant; detoxification; metal compound; soil; sediment; primer; aquatic environment; bioremediation; contamination; PCR; amplification.
                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a PCR mutagenesis primer used to mutagenise the marA coding sequence derived from transposon Tn21, to adapt it for plant expressibility. The nucleic acid molecule produced is useful for producing transgenic plants that are resistant to toxic heavy metals (especially mercury) and so can be used remediate and/or revegetate contaminated areas. The bacterial merA gene is derived from the transposon Tn21, which was originally isolated from the Incompatibility Group IncFII resistance plasmid NR1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Organometal; resistance; transcription; translation; regulation; merA;
                                                                                                                                                                                                                                                                                                            DNA encoding mercuric ion reductase in plant-expressible form – for producing transgenic plants resistant to toxic heavy metals {\sf res}
Metal ion resistance; mercuric ion reductase; merA; mercury; toxic heavy metal; transgenic plant; mutagenic; bacterial; transposon; Tn21; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   36
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00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length:
Matches:
Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 36 BP; 8 A; 13 C; 9 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-375-514-22 (1-205) x AAT92245 (1-36)
                                                                                                                                                                                                                                                                                                                                                         Example 1; Column 45; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  75 ProAlaAlaProGlyAlaAlaAlaGly 83
                                                                                                                                                                                                                         (UYGE-) UNIV GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   35 CCTGCTGCGCCGGGTGAGGCTGCAGGA 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer 3'N for Tn21 merA gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7.31e+04
41.00
88.89%
88.89%
3.71%
                                                                                                                                                               95US-0427097
                                                                                                                                                                                           95US-0427097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ28595/c
ID AAZ28595 standard; DNA; 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                      Summers AO;
                                                                                                                                                                                                                                                                                  WPI; 1997-470112/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity:
Query Match:
                                                                       Transposon In21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ransposon Tn21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Percent Similarity:
                                                                                                                                                               21-APR-1995;
                                                                                                                                                                                           21-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-DEC-1999
                                                                                                    JS5668294-A.
                                                                                                                                                                                                                                                      Meagher RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USS965796-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-0CT-1999.
                                                                                                                                  16-SEP-1997
                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ28595;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 31
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Sun Jun

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The invention relates to a nucleic acid molecule comprising a coding sequence for an organometal resistance protein, which is operably linked downstream of and under the regulatory control of a plant-expressible transcription and translation regulatory sequence. The organometal resistance gene is especially the merA gene from the transposon Th21 or the E.coli merB gene. merA encodes a mercuric ion reductase and merB encodes a corganomercury lyase. The nucleic acid sequences are useful for producing transgenic plants which are capable of efficiently processing mercury. These plants are able to remove or detoxify metal compounds such as methyl mercury and ionic mercury from soil, sediment, and aquatic environments providing an efficient bioremediation of metal and organometal contaminated sites.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This sequence represents a PCR primer used to amplify the Tn21 merA gene which is mutated by random mutagenesis (see AAZ28584-Z28589 for mutated sequences) and the mutated sequences are expressed in the
                                                                                                                                                                                                                     Metal resistance sequences for producing transgenic plants
                                                                                                                                                                                                                                                               Example 3; Column 17-18; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transgenic plants of the invention.
                                                                                     (UYGE-) UNIV GEORGIA RES FOUND INC.
                                                                                                                                Summers AO;
97US-0878957.
                                         95US-0427097
                                                                                                                                Meagher RB,
                                                                                                                                                                          WPI; 1999-579950/49
19-JUN-1997;
                                           21-APR-1995;
                                                                                                                              Rugh CL,
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Sequence 36 BP; 8 A; 13 C; 9 G; 6 T; 0 other;

36 0 0 0 0 Matches: Conservative: Mismatches: Indels: 7.31e+04 41.00 88.89% 88.89% 3.71% Similarity: Percent Similarity: Alignment Scores: Best Local S Query Match:

US-09-375-514-22 (1-205) x AAZ28595 (1-36)

75 ProAlaAlaProGlyAlaAlaAlaGly 83 35 ccrecrececedergageracage 9 ò g

AAQ41913 standard; DNA; 38 BP AAQ41913;

30-SEP-1993 (first entry)

Purine; promoter; human; erb B2/neu; gene; HER-2; homologue; rat; neu; triplex-forming oligonuclectide; TFO; retinoic acid; transgenic mice; core promoter element; growth factor; c-AMP; cancer; mammary tumour; tumour; NIH3T3 cells; pyrimidine; TPA; major groove; target; CAT box; TATA box; transcription; transforming; AT box protein; RNA polymerase; TFIId; control isomer; expression; ss.

erb B2/neu promoter randomised isomer, control 2.

Synthetic.

WO9309788-A.

27-MAY-1993

92WO-US09202. 28-OCT-1992; 91US-0792319. 13-NOV-1991;

(BAYU) BAYLOR COLLEGE MEDICINE

Hogan ME;

WPI; 1993-182231/22.

Use of triplex-forming oligo-nucleotide - to inhibit proliferation of cells contg. an erb. B2/neu gene site, for treating cancers, psoriasis etc.

Disclosure; Page 10; 26pp; English.

The sequences given in AAQ41911-13 are control isomers which comprise randomised sequences, based on triplex-forming isomers (TFO), which do not bind to the erb B2/neu target sequence, and have no effect on orb bayneu expression. The erb B2/neu (HER-2) gene is the human homologue of the rat neu gene. This human homologue is frequently amplified in tumours. When expressed at high levels in NHB73 cells, erb B2/neu is strongly transforming and results in a high incidence of mammary tumours in transgenic mice. The core promoter element of the B2/neu resides within a 300 bp region of the 5' flanking domain. This region contains elements which confer sensitivity to enhance core. The E2/neu resides within a 300 bp region of the 5' flanking domain. The promoter function in the presence of cell growth factors such as TPA, core and retinoic acid. Therefore, overexpression of of the B2/neu may be one mechanism leading to cancer initiation or expression. The sequences given in AAQ41905-10 are TFOS which are specific to the promoter region of erb B2/neu. They bind to the major groove of the complex to form a triplex. The TFOS are complementary location in the DNA duplex is a GC pair and T when the complementary location in the DNA duplex is an AT base pair. The target sequence such they include a G when the complementary location in the DNA duplex is an AT base pair. The target side for these TFOS should have a stretch of DNA which is at least 65% puring or these TFOS should have a stretch of DNA which is at least 65% puring or the CAT box and the TATA box. Inhibition at the CAT box and the TATA box. Inhibition of the erb protein with TFIId at the TATA box. Inhibition of the erb procession of the gene and may therefore be used to treat or prevent cancers.

Sequence 38 BP; 0 A; 0 C; 29 G; 9 T; 0 other;

1 6 2 0 8 38 Conservative: Mismatches: Matches: Indels: Length: Gaps: 7.72e+04 41.00 50.00% 50.00% 3.71% Best Local Similarity: Query Match: Percent Similarity Alignment Scores: Pred. No.: Score:

US-09-375-514-22 (1-205) x AAQ41913 (1-38)

44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro 59 æ g

AAQ44773/c RESULT 33

AAQ44773 standard; DNA; 38

AAQ44773;

(first entry) 12-0CT-1994

Steroid hormone related control oligonucleotide HIV CTR.

Triplex; triple helix; steroid hormone dependent transcription; regulation; progesterone receptor; HIV CTR; control; ss.

Synthetic. 10-NOV-1994

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/note= "Propanolamine group attached to this base."
                                                                                                                                                                                                                                                                                                                                                                                                                                                             directed against the progesterone receptor gene and inhibits progesterone-dependent transcription. The region of the target protected from DNaseI digestion by the PREZAP TVO overlaps more than one half-site of each progesterone receptor element (PRE). Two other TFOs were designed as controls of PREZAP activity; PRESP (AAA4472) has the PREZAP sequence inverted in a strand of opposite polarity and "HIV CTR" (AAA44773) contained an unrelated sequence of G and T residues with similar length and G/T ratio. Unlike with PREZAP, no triplexes were formed with either HIV CTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            therapy; infection; herpes simplex virus; human papilloma virus;
Epstein-Barr virus; HIV, adenovirus; respiratory syncitial virus;
hepatitis B virus; human cytomegalovirus; ss.
                                                                                                                                                                                                                                                                                                       Triple-helix forming oligo:nucleotide(s) to steroid hormone receptor target - useful for inhibiting steroid hormone induced gene activation and receptor transcription
                                                                                                                                                                                                                                                                                                                                                                                                                                        The triplex forming oligonucleotide (TFO) PRE2ap (AAQ44771) is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Guanosine; tetrad; inhibition; replication; virus; treatment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Guanosine rich oligonucleotide used to treat viral infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length:
Matches:
Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 38 BP; 0 A; 0 C; 29 G; 9 T; 0 other;
                                                                                                                                                                                                                         Tsai M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-375-514-22 (1-205) x AAQ44773 (1-38)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                               Example 5; Page 25; 47pp; English.
                                                                                                                                                                                                                      Hogan ME, Ing NH, Omalley BW,
                                                                                                                                                                          (BAYU ) BAYLOR COLLEGE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ79202 standard; DNA; 38 BP
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41.00
50.00%
50.00%
3.71%
                                                                                       93WO-US08329.
                                                                                                                               92US-0941362
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                                                                                                                                                                                                                                                                  WPI; 1994-100812/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Percent Similarity:
                                                                                     02-SEP-1993;
                                                                                                                               04-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alignment Scores:
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WO9405268-A.
                                       17-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9425037-A
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NAME OF THE PROPERTY OF THE PR
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The oligonucleotides (See AAQ79201-52) can be used to treat viral infections. The oligonucleotides inhibit viral replication by forming guanosine tetrads which form a stabilised 3D structure. Preferred oligonucleotides contain at least 2 runs of at least 2 guanosine bases and may be capped at the 3' terminus with a modifier selected from polyamine, poly-L-lysine, cholesterol and propanolamine. They may also have a modified phosphodiester linkage or be modified to contain a phosphorothioate linkage. They are used to treat infections with viruses such as herpes simplex virus, human papilloma virus, Epstein-Barr virus, HIV, adenovirus, respiratory syncitial virus, hepatitis B virus or human cytomegalovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human immunodeficiency virus; HIV; Epstein Barr virus; EBV; herpes simplex virus; HSV; human papilloma virus; HPV; adenovirus; respiratory syncytial virus; RSV; cytomegalovirus; CWV; hepatitis B; integrase inhibition; guanosine tetrad; ss.
                                                                                                                                                                                                                                                                                                     Oligo-nucleotide(s) rich in guanosine which form guanosine tetrads - used to treat viral infections, e.g. herpes-virus and
                                                                                                                                                                                                        Zendegui JG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  32 CCCACCCCA------CCCCCCCCACACCCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  38
0
1
1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length:
Matches:
Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                        Ojwang JO, Rando RF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 38 BP; 0 A; 0 C; 29 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Viral integrase inhibiting oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-375-514-22 (1-205) x AAQ79202 (1-38)
                                                                                                                                                                                                                                                                                                                                                                                                             Claim 41; Page 45; 101pp; English.
                                                                                                                          COLLEGE MEDICINE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT51620 standard; DNA; 38 BP.
                                             93US-0053027.
94WO-US04529.
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41.00
50.00%
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3.71%
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                                                                                                                          (BAYU ) BAYLOR COLLEGE MED (TRIP-) TRIPLEX PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-NOV-1997 (first entry)
                                                                                                                                                                                                        Hogan ME,
                                                                                                                                                                                                                                                        WPI; 1994-357890/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                        Pennewald S,
25-APR-1994;
                                                                          28-OCT-1993;
                                                23-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9703997-A1
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19-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-JUL-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAT51620;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match:
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columnar box structure"

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Query Match:
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AAQ95496/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              δŽ
Column; box; stacked tetrad; inhibition; replication; pathophysiological; herpes simplex virus; HSV; human papilloma virus; Epstein Barr Virus; HPV; EBV; HIV; human immunodeficiency virus; adenovirus; RSV; HBV; HCMV; respiratory syncytial virus; hepatitis B virus; human r-cell leukaemia virus; HTV; ss.
                                                                                                                                                                                                                                                                                        AMT51619-T51698 are oligonucleotides used to inhibit the production of viruses within a host cell. The oligonucleotides may form guanosine tetrads (structures formed of eight hydrogen bonds by coordination of the four oxygen atoms of guanine with alkali cations believed to bind to the centre of a quadruplex, and by strong stacking interactions) and are used to prevent the integration of viral nucleic acid into a host genome. The oligonucleotides inhibit functioning of the integrase enzyme and hence prevent viral infection. Viral infections that may be treated include human immunodeficiency virus (HIV), Epstein Barr virus (EBV), herpes simplex virus (HSV), human papilloma virus (HPV), adenovirus, respiratory syncytial virus (RSV), cycomegalovirus (CMV) and hepatitis B virus (HBV), especially HIV-1 infection.
                                                                                                                                                                                              Oligo:nucleotide(s) capable of forming guanosine tetrads - inhibit viral enzyme responsible for integrating viral nucleic acid into the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide #4 forms an intramolecular stacked tetrad structure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "forms intramolecular stacked tetrad or 3D
                                                                                                                  Pommier Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "optionally contains phosphodiester internucleotide linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               44 ProAlaProGlyIlePheSerSerGlnProGlyHisThrProHisPro 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1070838
                                                                                                                  Hogan ME, Mazumder A, Ojwang JO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 38 BP; 0 A; 0 C; 29 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-375-514-22 (1-205) x AAT51620 (1-38)
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1..38
/*tag= a
                                                                                                                                                                                                                                                                 Claim 3; Page 141; 245pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX79211 standard; DNA; 38 BP
 95US-0535168.
96US-0013688.
96US-0014007.
96US-0015714.
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                                                                                    (ARON-) ARONEX PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag=
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                                                                                                                                     Rando RF, Zendegui JG;
                                                                                                                                                                WPI; 1997-132569/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               32 CCCACCCCA----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   misc_structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Percent Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Key
modified_site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alignment Scores:
                                  25-MAR-1996;
17-APR-1996;
                    19-MAR-1996;
                                                                                                                     Fennewald S,
     23-OCT-1995;
                                                                                                                                                                                                                                    host genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match:
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Sequences AAX79210-X79275 represent oligonucleotides (ON) which are able to form a columnar box or "stacked tetrad" structure by intramolecular internucleotide binding. The ONS are used to inhibit the replication of viruses. They are able to suppress virus production for prolonged periods after an initial short treatment regimen. They can be used for treating pathophysiological states caused by viruses such as herpes simplex virus, human papilloma virus, Epstein Barr Virus, Human average of the periods of the p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cdn-1; apoptosis modulator; adoptive immunotherapy; therapy; HIV; autoimmune disease; reperfusion injury; hepatitis, osteoporosis; shock; lymphoma; eczema; primer; polymerase chain reaction; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New guanosine-rich tetrad forming oligonucleotide(s) - used for inhibiting virus replication for treating e.g. herpes simplex, papilloma, HIV, adenovirus or hepatitis B virus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       44 ProAlaProGly1lePheSerSerGlnProGlyHisThrProHisPro 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wallace TL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -------ccccccacaccccacacccc 3
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Mismatches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ojwang JO, Rando RF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length:
Matches:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    adenovirus, respiratory syncytial v
cytomegalovirus and HTLV I and II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВÞ
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/mod_base= i
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                                                                                                                                                                  98WO-US01974
                                                                                                                                                                                                                                                                                               97US-0037374
                                                                                                                                                                                                                                                          97US-0987574
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                   (ARON-) ARONEX PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        32 cccacccca-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cossum PA, Hogan ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cdn primer Bclx 2-39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-446809/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Alignment Scores:
WO9833807-A1
                                                                                                                                                                       )3-FEB-1998;
                                                                                                                                                                                                                                                          09-DEC-1997;
                                                                                                                                                                                                                                                                                               04-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-NOV-1995
                                                                                   06-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ95496;
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New nucleic acid sequences encoding Cdn apoptosis modulators - and related vectors, transformed cells, proteins and antibodies, useful or diagnosis and treatment e.g. of HIV infection, reperfusion injury
                                                                                                                                                                                                                                                                                                                    In order to identify new Bcl-2 family members, degenerate PCR primers (AAQ95495-96) based on sequences in these regions were used in the amplification of human heart and B lymphoblastoid cell line WI-L2 cDNA. Cdn-1 cDNA (AAQ95492) was obtd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; detection of early stage allergic disease; atopic dermatitis; antiallergic; eosinocyte; eosinophil; bcl-2; probe; ss.
             /*tag= c
/note= "base n at position 25 is not identified"
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/note= "base n at position 28 is not identified"
                                                                                                                                                                                                                                                                                                                                                                                                             00700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag= a
/mod_base= OTHER
/note= "OTHER= labelled with FAM
(6-carboxy-fluorescein)"
                                                                                                                                                                                                                                                                                                                                                                                                            Length:
Matches:
Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fluorescent probe for DNA encoding human bcl-2.
                                                                                                                                                                                                                                                                                                                                                                           Sequence 39 BP; 7 A; 12 C; 5 G; 9 T; 6 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-375-514-22 (1-205) x AAQ95496 (1-39)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure, Fig.1; 66pp; English.
                                                                                                                                                                           (LXRB-) LXR BIOTECHNOLOGY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK52487 standard; DNA; 24 BP
                                                                                                                                                       93US-0160067.
                                                                                                                                                                                                                                                                                                                                                                                                         7.93e+04
41.00
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                                                                                                                       94WO-US13930.
                                                                                                                                           94US-0320157
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             /*tag=
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/*tag=
                                                                                                                                                                                                 Barr PJ, Kiefer MC;
                                                                                                                                                                                                                     WPI; 1995-215106/28
 misc_difference 25
                                misc_difference 28
                                                                                                                                                                                                                                                                                                                                                                                                                              Percent Similarity:
Best Local Similarity:
Query Match:
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modified_base
                                                                                                                     30-NOV-1994;
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                                                                                                                                          07-OCT-1994;
                                                                                                                                                      30-NOV-1993;
                                                                           WO9515084-A
                                                                                                                                                                                                                                                                                                                                                                                               Alignment Scores:
                                                                                               08-JUN-1995
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                                                                                                                                                                                                                                                                                                                                                                                                          No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BK52487
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The present invention relates to a method for detecting early stage allergic diseases, particularly atopic dermatitis. The method comprises determining the expression levels of granulocyte macrophage colony stimulating factor receptor alpha or beta (GM-CSF R-alpha or beta), interleukin 3 receptor alpha (IL-3 R-alpha), bcl-2, bcl-x, platelet activation factor receptor (PAF R) or CD44 in eosinocytes of a subject to be tested. The method further comprarison with expression levels in healthy volunteers. The method is useful for the early diagnosis and treatment of early stage allergic diseases such as atopic dermatitis. The present sequence represents a probe used in the methods of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cytokine; tumour necrosis factor-alpha; TNF-alpha; lipopolysaccharide; mitogen-induced; colinear triplex; capping group; cholesterol; tumour cell; glioblastoma cell; ss.
                                                                                                                                                                                                                                                    Detecting early stage allergic diseases with markers of 7 genes of GM-CSF R-beta, GM-CSF R-alpha, IL-3 R-alpha, PAF R, bcl-2, bcl-x and
              /note= "OTHER= labelled with TAMRA
   (6-carboxy-N,N,',N'-tetramethylrhodamine)"
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Mismatches:
Indels:
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/note= "linked to cholesterol"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytokine production inhibitor J111-56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-375-514-22 (1-205) x ABK52487 (1-24)
                                                                                                                                                                           (GENO-) GENOX SOYAKU KENKYUSHO KK. (KOKU-) KOKURITSU SHONI BYOIN INCHO.
                                                                                                                                                                                                                                                                                                                Example 1; Page 20; 25pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
   /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 143 AsnTrpGlyArgIleValAla 149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP.
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                                                                                                                                               11-OCT-2000; 2000JP-0311193.
                                                                                                                   11-OCT-2000; 2000JP-0311193
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                                                                                                                                                                                                                       WPI; 2002-475327/51.
                                                                                                                                                                                                                                                                                  CD44 in eosinophils
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Percent Similarity:
Best Local Similarity:
Query Match:
                                                        JP2002119281-A.
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modified_base
                                                                                       23-APR-2002.
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Rando RF,

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human vWF positive sequences using two synthetic DNA probes (AAQ13426-7). The probes were synthesised according to the published DNA sequence (Sadler et al., Proc. Nat. Acad. Sci. 82: 6394-8 (1985) and Verweij et al., EMBO J. 3: 1829-47 (1986)) of human VWF.
                                                                                                                                                                     The invention relates to an apoptosis inhibitor comprising the amino acid sequence of Bc1-2 protein in which at least one serine residue is substituted by alanine or aspartic acid. The protein has increased apptosis inhibitory activity compared with the wild type Bc1-2 protein. The mutated Bc1-2 protein is useful in the treatment of disorders caused by apoptosis. The present sequence was used to create a mutant Bc1-2 protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A human endothelial cDNA library in lambda gt11 was screened for
                                                                                   New apoptosis inhibitors, useful for treating apoptosis related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Non-glycosylated peptide(s) of vWF GPlb binding domain inhibits platelet adhesion and aggregation for treatment of cerebrovascular and cardiovascular disorders
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                                                                                                                                                                                                                                                                                                                              Sequence 27 BP; 2 A; 9 C; 12 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                      Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            vWF GPIb binding domain probe (4700-4739).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-375-514-22 (1-205) x AAH45300 (1-27)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 26; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27 GTCGCCAGGACCGCGCCGCTGCAGACC 1
                                                                                                                                           Example 1; Page 10; 43pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ13427 standard; DNA; 40 BP
                                                                                                                                                                                                                                                                                                                                                                                    6.34e+04
40.00
100.00%
88.89%
3.62%
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Garfinkel L, Richter T;
             Shibazaki F, Kuma H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1991-281423/38.
                                                 WPI; 2001-381681/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                              Percent Similarity:
Best Local Similari
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          32-MAR-1990;
                                                                                                                                                                                                                                                                                                                                                                        Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ13427;
                                                                                                             disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match:
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No.:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dp
AAT08787-T08792 represent oligonucleotides capable of inhibiting cytokine production. These sequences specifically inhibit tumnour necrosis factor-alpha (TNF-alpha) production. They also inhibit tumnour lipopolysaccharide or other mitogen-induced production of TNF-alpha. The sequences were designed to form a colinear triplex with a region of the TNF-alpha gene. The triplex is stabilised by the guanosine tetrads, or the runs of at least 2 contiguous quanosines. This sequence targets the region from 1429 to 1456 of the TNF-alpha gene (relative to the mRNA start site). All of the sequences have a 3' capping group (either an anne or a cholesterol) to confer stability against nuclease digestion. The sequences can be used to inhibit the growth of tumour cells, particularly TNF-alpha responsive tumour cells (preferrably glioblastoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; Bc1-2; gene therapy; apoptosis inhibitor; mutant; primer; ss
                                                                                                                                                                                                                                           Guanosine-rich oligo:nucleotide cpds. - useful for inhibiting cytokine prodn., TNF-alpha prodn. and the growth of TNF-alpha responsive cells, e.g. tumour cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Bcl-2 mutagenic primer oligo-3 for S70A substitution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     00000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 27 BP; 0 A; 0 C; 18 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (1-27)
                                                                                                                                                                                                                                                                                                                      Claim 14; Page 17; 29pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HISM ) HISAMITSU PHARM CO LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-375-514-22 (1-205) x AAT08792
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH45300 standard; DNA; 27 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            53 ProGlyHisThrProHisPro 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21 CCCACACACACCCCCACCCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99JP-0350427.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-DEC-2000; 2000WO-JP08667.
                                                       95WO-US06987
                                                                                             94US-0250862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       34e+04
                                                                                                                                   (TRIP-) TRIPLEX PHARM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       40.00
85.71%
85.71%
3.62%
                                                                                                                                                                         Santon JB;
                                                                                                                                                                                                           WPI; 1996-030253/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-SEP-2001
                                                         31-MAY-1995;
                                                                                             31-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Alignment Scores:
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                     07-DEC-1995
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AAH45300;

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RESULT 40 AAH45300/

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Query Match:

cells).

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This synthetic oligonucleotide corresponds to nucleotides flanking the 3' end of the human von Willebrand Factor (vWF) GPIb binding domain coding region. It was used as a probe for the detection of vWF sequences in a human endothelial cell cDNA library. A GPIb binding domain 5' probe was also used (see AAV74158). A series of vWF cDNA clones covering the entire GPIb binding domain (see AAV74157) were identified and isolated. The invention provides non-glycosylated, biologically active polypeptides (see AAW90119-20) which comprise the vWF GPIb binding domain. These are used in claimed methods for treating and preventing cerebrovascular and cardiovascular disorders, and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Composition for inhibiting blood platelet aggregation - containing polypeptide comprising von Willebrand factor GPlb binding domain
                                                                                                                                                                                                                                                                                                                                                                                                von Willebrand Factor; vWF; human; GPIb binding domain; antiaggregant; platelet aggregation; cerebrovascular disorder; cardiovascular disorder; myocardial infarction; angina; thrombolytic; platelet adhesion; therapy; probe; ss.
                                                                         0 0 0 0 0 0
                                                                                                  Conservative:
Mismatches:
Indels:
                          Sequence 40 BP; 11 A; 14 C; 8 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                       Human mature von Willebrand Factor DNA probe.
                                                                       Length:
Matches:
                                                                                                                                              Gaps:
                                                                                                                                                                   US-09-375-514-22 (1-205) x. AAQ13427 (1-40)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BIOT-) BIO-TECHNOLOGY GENERAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Column 12; 50pp; English.
See also AAQ13426-28 and AAR13939.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inhibiting platelet aggregation.
                                                                                                                                                                                                  53 ProGlyHisThrProHisPro 59
                                                                                                                                                                                                                   1 CCAGGACGAACGCCACATCCA 21
                                                                                                                                                                                                                                                                                    AAV74159 standard; DNA; 40 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 91US-0753815.
90US-0487767.
91WO-US01416.
93US-0080690.
                                                                    9.47e+04
40.00
85.71%
85.71%
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                                                                                                                                                                                                                                                                                                                                            12-APR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Garfinkel L, Richter T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-069781/06.
                                                                                                                 Similarity:
                                                                                              Percent Similarity:
Best Local Similarit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       domo sapiens.
                                                        Alignment Scores:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAR-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                 AAV74159;
                                                                                                                             Query Match:
                                                                                                                                                                                                                                                                       AAV74159
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This sequence represents a probe for DNA encoding the protein of the invention. The protein of the invention is the GPlb binding domain of the invention. The protein of the invention is the GPlb binding domain of the von Willebrand factor. The protein is used for inhibiting platelet aggregation, especially for treating cerebrovascular disorders. It is also used for treating cardiovascular disorders, especially acute myocardial infarction or angina. The protein is also used for inhibiting platelet aggregation before, during or after angioplasty, thrombolytic treatment or coronary bypass surgery, for maintaining blood vessel or coronary before, during or after coronary bypass surgery. It can also be used for inhibiting thrombosis, optionally associated with an inflammatory response, for inhibiting platelet adhesion to damaged constrains or devices, for inhibiting re-occlusion after angioplasty or thrombolysis, or for thrombolytic treatment of thrombi-containing platelet-rich aggregates.
                                                                                                                                                                                                                                                                                                                                                                                                                          Von Willebrand factor; GP1b binding domain; inhibitor; re-occlusion; platelet aggregation; cerebrovascular disorder; cardiovascular disorder; angloplasty; thrombi-containing platelet-rich aggregate; thrombosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polypeptide comprising von Willebrand factor GPlb binding domain
                                                 0 0 0 0 0 0
                                                Length:
Matches:
Conservative:
                                                                                                                                                                                                                                                                                                                                                                                           Probe for Von Willebrand factor coding sequence.
Sequence 40 BP; 11 A; 14 C; 8 G; 7 T; 0 other;
                                                                                                 Mismatches:
                                                                                                                   Indels:
                                                                                                                                Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful as platelet aggregation inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Column 13-14; 62pp; English.
                                                                                                                                                              US-09-375-514-22 (1-205) x AAV74159 (1-40)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BIOT-) BIO-TECHNOLOGY GENERAL CORP.
                                                                                                                                                                                                                   1 CCAGGACGAACGCCACATCCA 21
                                                                                                                                                                                                                                                                                              AAV08903 standard; DNA; 40 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      91WO-US01416.
90US-0487767.
91US-0753815.
93US-0080690.
                                                 9.47e+04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94US-0347594
                                                                                                                                                                                                53 ProGlyHisThrProHisPro
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                                                               40.00
85.71%
85.71%
3.62%
                                                                                                                                                                                                                                                                                                                                                            26-FEB-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              therapy; probe; ss.
                                                                             Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-MAR-1990;
03-SEP-1991;
22-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US5849536-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-DEC-1998.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                             AAV08903;
                                                                                                                   Query Match:
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0
0
                                                                                                                                                                                                                                                                 RESULT 43
                                                                                                                                                                                                                                                                                  AAV08903
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This is a probe for the human von Willebrand factor (VWF). The invention relates to methods of treating arteriosclerosis and stenosis using polypeptides derived from the human VWF GPIb binding domain. VWF is a large plasma protein which is synthesised in the endothelial cells of the blood vessels, and also by megakaryocytes which are the precursors of platelets. The treatment methods involve the administration of a polypeptide (see AAVG7242 or AAVG7243). These polypeptides contain a fragment of the VWF GPIb binding domain amino acid sequence. The polypeptides inhibit platelet adhesion and aggregation, therefore the methods may be used for preventing arteriosclerosis and stenosis (as a result of smooth muscie proliferation following vascular injury). The polypeptides may also be used for treating other cerebrovascular and
                                                                                                                                                                                                                                                                                                                                                                                         Von Willebrand Factor; vWF; GPIb binding domain; arteriosclerosis; platelet adhesion inhibitor; stenosis; cerebrovascular injury; human; thrombosis; cardiovascular injury; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Preventing arteriosclerosis and stenosis using polypeptides derived from the human von Willebrand factor GPIb binding domain -
                                                                                                                                                                                                                                                                                                                                                                Probe for human von Willebrand Factor (vWF) GPIb binding domain.
                                             0 0 0 0 0
                                           Length:
Matches:
Conservative:
Mismatches:
Indels:
T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 7 T; 0 other,
                                                                                                                     Gaps:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cardiovascular injuries and thrombosis
                                                                                                                                                US-09-375-514-22 (1-205) x AAV08903 (1-40)
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                                                                                                                                                                               53 ProGlyHisThrProHisPro 59
                                                                                                                                                                                                1 CCAGGACGAACGCCACATCCA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       91US-0753815.
93US-0080690.
90US-0487767.
                                        9.47e+04
40.00
85.71%
85.71%
3.62%
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 A; 14 C;
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                                                                                                                                                                                                                                                                       AAZS6179 standard; DNA; 40
                                                                                                                                                                                                                                                                                                                                 27-MAR-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Garfinkel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-096379/08.
                                                                         Percent Similarity:
Best Local Similarity:
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22-JUN-1993;
02-MAR-1990;
01-MAR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-JUN-1995;
                               Alignment Scores:
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 Sequence 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US6008193-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Richter T,
                                                                                                                                                                                                                                                                                                     AAZ56179;
                                                                                                     Query Match
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                                                                                                                                                                                                                                         RESULT 44
                                                                                                                                                                                                                                                       AA256179
                                                             Score:
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    A. fumigatus codon optimised phytase gene constructing phy52r PCR primer.

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                                                                                                                                                                                                                                                                                                                Phytase; Pichia cell; heterologous nucleotide; codon optimisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel Pichia cell useful for producing polypeptide of interest, comprises a copy of heterologous nucleotide sequence encoding polypeptide of interest
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 047140
  90100
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Matches:
Conservative:
                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches:
Indels:
                             Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 40 BP; 9 A; 5 C; 17 G; 9 T; 0 other;
  Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tsutsumi N, Taira R;
                                          Indels:
                                                                                  (1-40)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Page 53; 61pp; English.
                                                                                  US-09-375-514-22 (1-205) x AAZ56179
                                                                                                                                                                                                AAD17995 standard; DNA; 40 BP
                                                                                                              53 ProGlyHisThrProHisPro 59
                                                                                                                               1 CCAGGACGAACGCCACATCCA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9.47e+04
40.00
66.67%
58.33%
3.62%
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15-MAR-2000; 2000DK-0000419.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-MAR-2001; 2001WO-DK00154
                                                                                                                                                                                                                                                         (first entry)
40.00
85.71%
85.71%
3.62%
                                                                                                                                                                                                                                                                                                                                                        Aspergillus fumigatus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NOVO ) NOVOZYMES AS.
    Percent Similarity:
Best Local Similarity: 1
Query Match:
DB:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Terui Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-582273/65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                WO200166693-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Percent Similarity:
                                                                                                                                                                                                                                                        10-DEC-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                               13-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Takagi S,
                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention
                                                                                                                                                                                                                            AAD17995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match:
                                                                                                                                                                       RESULT 45
```

US-09-375-514-22 (1-205) x AAD17995 (1-40)

40

9.47e+04

Alignment Scores: Pred. No.:

Sun Jun

(first entry)

19-JUL-2000

AAA26253;

```
The present invention relates to a mouse model expressing a modified growth hormone and designated SMA1. The model is used to study diseases, test prospective treatments and identify early detection markers associated with abnormal growth hormone (GH) or ICF-1 activity. Particularly dwarfism, Laron syndrome, Illig-type deficiency, Kowarsky syndrome, pituitary gland defliciency, reduced bone mineral density, increased or insufficient body fat accumulation, diabetes, osteoporosis, glomerulosclerosis, gigantism, acromegaly, information processing and connection defects, age-related memory and behaviour deflicits, glucose metabolism disorders and cardiovascular disorders. The present sequence is a PCR primer described in the exemplification of the
                                                                                                                                                                                                                                            Mouse; human; animal model; growth deficiency; cognitive function; growth hormone; IGF-1, osteoporosis; cardiovascular disorder; dwarfism; Illig-type deficiency; Kowarsky syndrome; diabetes; acromegaly; gigantism; glomerulosclerosis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New animal model expressing a modified growth hormone has reduced body weight and size and is useful to determine treatment for growth hormone, particularly IGF-1 related conditions and diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ŝ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wolf
                                                                                                                                                                                                           Murine growth hormone coding sequence PCR primer SEQ ID NO: 16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Schlueter V, Wattler F, Wattler S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              470700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 40 BP; 5 A; 15 C; 16 G; 4 T; 0 other;
186 HisThrTrpIleGlnAspAsnGlyGlyTrpValGly 197
                                  2 CATGGTAGCGTTCAATGGGAAGGTGGCTGGGTTGGA 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-375-514-22 (1-205) x AAL46236 (1-40)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 33 GlyAspValGlyAlaAlaProProGly 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     40 GGAGACTCGGGAGCTGTGCTCCCGGC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 6; Page 74; 93pp; English.
                                                                                                        BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA26253/c
ID AAA26253 standard; DNA; 23 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9.47e+04
40.00
77.78%
77.78%
3.62%
                                                                                                                                                                                                                                                                                                                                                                                                                                  02-APR-2001; 2001WO-EP03733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAR-2000; 2000DE-1016083
                                                                                                      AAL46236 standard; DNA; 40
                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (INGE-) INGENIUM PHARM AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cornali E, Nehls MC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-616425/71.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Percent Similarity:
Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                 WO200172119-A2.
                                                                                                                                                                                                                                                                                                                                 Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alignment Scores:
                                                                                                                                                                           15-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                  04-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention.
                                                                                                                                         AAL46236;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No.:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 47
                                  q
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The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(d1)thioate with a target sequence and contain at least one phosphoro(d1)thioate link, having endomuclease activity. (A), and more generally any catalytic nucleic acid (A) that modulates expression of the coestrogen receptor gene, are used to treat cancer (particularly of breast or receptor gene, are used to treat cancer (particularly of breast or conformal or for other conditions associated with levels of coestrogen receptor. Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, particularly for identification of therapeutic targets, and as research reagents (for RNA, in the same way that restriction endonucleases are used with DNA). The combination of modifications in (A) improves resistance to nucleases, binding affinity and/or activity. AAA23503 to AAA24748 to AAA2592 represent their corresponding target sequences, and AAA26105 represent ceptor corresponding target sequences, and AAA26107 to AAA26211 represent their corresponding target sequences, and AAA2619 to AAA26211 represent chair corresponding target sequences. AAA26219 to AAA26211 represent content in the procession of the proce
                                                                                                                                                                                      hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bellon L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acids that interact, and optionally cleave, target
                                                                                                                                                               Jestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage;
                                                                                                                  3cl-2 antisense oligonucleotide sequence SEQ ID NO:2751.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Beigelman L, McSwiggen JA, Karpeisky A,
Zwick M, Jarvis T, Woolf T, Haeberli P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 23 BP; 9 A; 9 C; 4 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length:
Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (1-23)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 9; Page 106; 148pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequences, used to treat cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-375-514-22 (1-205) x AAA26253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               149 AlaPhePheGluPheGlyGly 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GCCTTCTTTGAGTTCGGTGGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-US08547.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98US-0082404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    98US-0103636.
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39.00
100.00%
100.00%
3.53%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-013248/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Reynolds M, Zwic
Matulic-Adamic J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Percent Similarity:
                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thompson JD,
                                                                                                                                                                                                                                                                                                                                    W09954459-A2
                                                                                                                                                                                                                                                                                                                                                                                                                              19-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Alignment Scores:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match:
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δ
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21 GCCTTCTTTGAGTTCGGTGGG 1

AAH45301/c ID AAH45301 standard; DNA; 27

RESULT 49

AAH45301;

Human; Bcl-2; gene therapy; apoptosis inhibitor; mutant; primer; ss.

Human Bcl-2 mutagenic primer oligo-4 for S70D substitution.

10-SEP-2001 (first entry)

```
The present invention describes nucleic acids (A) that interact stably with a target sequence and contain at least one phosphoro(di)thioate clink, having endonuclease activity. (A), and more generally any catalytic nucleic acid (A') that modulates expression of the oestrogen catalytic nucleic acid (A') that modulates expression of the oestrogen receptor gene, are used to treat cancer (particularly of breast or condenderium), in vivo or by transforming cells at vivo and implanting cells or for other conditions associated with levels of catagets. Bob of the control of correlate inhibition of gene expression with cargets, and as research reagents (for RNA, in the same way that modifications in phenotype, particularly for identification of therapeutic restriction endonucleases are used with DNA). The combination of and/or activity. AAA23503 to AAA2447 represent oestrogen receptor corresponding target sequences, and AAA25992 represent their corresponding target sequences. AAA2593 to AAA26107 represent cerponding target sequences, and AAA26107 to AAA2618 represent corresponding target sequences, and AAA26107 to AAA2618 represent corresponding target sequences. AAA2619 to AAA2611 represent cotheir ribozyme sequences and antisense oligonucleotides used in the
                                                                                                                                                                               Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease; anticancer; breast cancer; endometrium cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Karpeisky A, Bellon L;
Haeberli P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acids that interact, and optionally cleave, target
                                                                                                                                             Bcl-2 antisense oligonucleotide sequence SEQ ID NO:2753.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                n L, McSwiggen JA,
Jarvis T, Woolf T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23 BP; 9 A; 9 C; 4 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 9; Page 106; 148pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequences, used to treat cancer
                                 AAA26255 standard; DNA; 23 BP.
                                                                                                                                                                                                                                                                                                                                                                                                99WO-US08547.
                                                                                                                                                                                                                                                                                                                                                                                                                                  98US-0082404.
98US-0103636.
                                                                                                         19-JUL-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thompson JD, Beigelman L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zwick M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-013248/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matulic-Adamic J;
                                                                                                                                                                                                                                                                                                                WO9954459-A2
                                                                                                                                                                                                                                                                                                                                                                                          19-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                      28-OCT-1999,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reynolds M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Alignment Scores:
                                                                       AAA26255;
RESULT 48
                   AAA26255,
```

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Human; Bcl-2; gene therapy; apoptosis inhibitor; mutant; primer; ss.
                                                                                                                                                                                                                                      Human Bcl-2 mutagenic primer oligo-6 for S116D substitution.
                  27
8
0
1
0
0
                                     Conservative:
                                                Mismatches:
                 Length:
Matches:
                                                         Indels:
                                                                                   US-09-375-514-22 (1-205) x AAH45301 (1-27)
                                                                                                         66 ValAlaArgThrSerProLeuGlnThr 74
                                                                                                                    27 GTCGCCAGGACCGACCGCTGCAGACC 1
                                                                                                                                                                      AAH45303 standard; DNA; 27 BP.
           7.38e+04
39.00
88.89%
88.89%
                                                                                                                                                                                                                 10-SEP-2001 (first entry)
                                             Similarity:
                             Percent Similarity:
Alignment Scores:
                                                                                                                                                                                            AAH45303;
                                                      Query Match:
                                            Best Local
                                                                                                                                                              AAH45303/c
                                                                                                                                                    RESULT 50
```

23 0 0 0

Conservative: Mismatches:

100.00% 100.00% 3.53%

Best Local Similarity: Percent Similarity:

Query Match:

Indels:

US-09-375-514-22 (1-205) x AAA26255 (1-23)

149 AlaPhePheGluPheGlyGly 155

Matches:

Length:

6.27e+04 39.00

amino acid sequence of Bc1-2 protein in which at least one serine residue is substituted by alanine or aspartic acid. The protein has increased apoptosis inhibitory activity compared with the wild type Bc1-2 protein. The mutated Bc1-2 protein is useful in the treatment of disorders caused by apoptosis. The present sequence was used to create a mutant Bc1-2 protein of the invention.

Sequence 27 BP; 2 A; 8 C; 12 G; 5 T; 0 other;

The invention relates to an apoptosis inhibitor comprising the

Example 1; Page 10; 43pp; Japanese.

disorders

New apoptosis inhibitors, useful for treating apoptosis related

(HISM) HISAMITSU PHARM CO LTD.

Shibazaki F, Kuma H; WPI; 2001-381681/40.

99JP-0350427.

09-DEC-1999;

07-DEC-2000; 2000WO-JP08667.

WO200142459-A1.

14-JUN-2001,

Homo sapiens.

Synthetic.

```
The invention relates to an apoptosis inhibitor comprising the amino acid sequence of Bc1-2 protein in which at least one serine residue is substituted by alanine or aspartic acid. The protein has increased apoptosis inhibitory activity compared with the wild type Bc1-2 protein. The mutated Bc1-2 protein is useful in the treatment of disorders caused by apoptosis. The present sequence was used to create a mutant Bc1-2 protein of the invention.
                                                                                                                                                                                                          New apoptosis inhibitors, useful for treating apoptosis related
                                                                                                                                                                                                                                                                                                                                                                                                                   27
0
0
0
0
                                                                                                                                                                                                                                                                                                                                                                           Sequence 27 BP; 5 A; 8 C; 9 G; 5 T; 0 other;
                                                                                                                                                                                                                                                    Example 1; Page 11; 43pp; Japanese.
                                                                                                                                     (HISM ) HISAMITSU PHARM CO LTD.
                                                                                      07-DEC-2000; 2000WO-JP08667.
                                                                                                               99JP-0350427.
                                                                                                                                                                 Shibazaki F, Kuma H;
                                                                                                                                                                                          WPI; 2001-381681/40.
                                       WO200142459-A1.
Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                            Alignment Scores:
                                                                                                                09-DEC-1999;
                                                               14-JUN-2001.
                                                                                                                                                                                                                                  disorders
```

Length:
Matches:
Conservative:
Mismatches:
Indels: Gaps: US-09-375-514-22 (1-205) x AAH45303 (1-27) 7.38e+04 39.00 88.89% 88.89% 3.53% Pred. No.: Score: Secore: Percent Similarity: Best Local Similarity: Query Match: ò

Search completed: May 31, 2003, 22:29:38 Job time : 222 secs

g

GenCore version 5.1.5 Copyright (c) 1993 - 2003 Compugen Ltd.

OM nucleic - nucleic search, using sw model

Run on:

Perfect score: Sequence:

US-09-375-514-19 5086 1 gcgcccgccctccgcgccg......caatgaatgtatataaaagc 5086

IDENTITY_NUC Gapop 10.0 , Gapext 1.0 Scoring table:

2054640 segs, 14551402878 residues earched:

770742 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 10 Maximum DB seq length: 40

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 500 summaries

GenEmbl:*

Database :

gb_sy:*
gb_un:*
gb_vi:*
em_ba:*
em_hun:*
em_hun:*
em_nu:*
em_nu:*
em_om:*
em_om:*
em_ow:*
em_ow:* gb_ba:*
gb_htg:*
gb_in:*
gb_om:*
gb_om:*
gb_pat:*
gb_ph:*
gb_ph:*
gb_pl:*
gb_ro:*
gb_ro:*

em_pl:* em_ph:

em_vi:*
em_htg_hum:*
em_htg_inv:*
em_htg_other:* em_htg_mus:*
em_htg_pln:*
em_htg_rod:* em_sts:* em_un:* em_ro:

em_htgo_mus:*
em_htgo_other:* em_htg_mam:* em_htg_vrt:* em_sv:* em_htgo_hum:*

Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

S	Description	nen	7288 Sequence	7550 Sequenc	7301 S	2604 Sequence	183 Sequence	7307 Sequenc	364 Sequence	7291 Sequenc	3626	340 Sequence	96235 Sequence	203	7294 Sequence	341 S	6236 Sequence	6240 Sequenc	M 603	91 Sequence 4	53 MBR repo 7303 Seguen	103 Sequence 2	123 Sequence	103 Method for	98 Sequence 2	.47 Sequence 3	06 Sequence 2	98 Sequence 1	68 Sequence 1	10 Sequence 1	Sequenc	77 Sequence 1 92 Sequence 1	46 Sequence 3	28 Sequence I	56 Sequence 6	86 Sequence 1	74 Sequence 8	25 Sequence	0/ Sequence 1 88 Sequence 4	7731 Sequence	40/	1257 Sequence	86 Sequence 2	os sequence 3 26 Sequence 2	61 Sequence 3	13	* 1011111111111111111111111111111111111
SUMMARIE	ΙD	AX377551	28	55	2 6	60	33	AR00/302 AR007307	4	29	AR096260	90	AR096235	ğ ლ	AR007294	A56841	AR096236	24	E32209 I31263	131491	A22453 AR007303		I31323 T31359	E32203	131298	131464 131147	131306	131260	131268	131280 131410	131165	1312// 131192	131446	131228 AX028364	9	131286 131485	131174	I31225	131488	AX477731	~ 0	AR051257	I28386	131326	131461	I31419 T31494	* ` * * * * *
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d	Query Match	0.8																																													
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Mammalia; Eutheria; Primates; Catarrhini; Homfnidae; Homo.
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Mammalia; Eutheria; Primates; Catarthini; Hqminidae; Homo.
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Nanogen Recognomics GmbH (DE)
Location/Qualifiers
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Muth.J.
Method for detecting mutations in nucleofide sequences
Patent: W0 0212553-A 29 14-FEB-2002;
Nanogen Recognomics GmbH (DE)
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Thompson, J.D. and Draper, K.G.
Method and reagent for treatment of diseases caused by expression
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Mammalia; Eutheria, Primates; Catarrhini; Hominidae; Homo.
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Method for detecting mutations in nucleotide sequences
Patent: WO 0212553-A 27 14-FEB-2002;
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Patent: US 5750390-A 15 12-MAY-1998;
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37; Conserwative 0; Mismatches
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0.7%; Score 37; DB
Best Local Similarity 100.0%; Pred. No. 05
Matches 37; Conservative 0; Mismatches
                                                                                                                                              of the bcl-2 gene
Patent: US 5750390-A 2/12-MAY-1998;
Location/Oyalifiers
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US 5750390.
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Sequence 27 from Patent W00212553.
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/db_xref="taxon:9606"
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AR007288.1 GI:3966772
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Thompson,J.D. and Draper,K.G.
Thompson,J.D. and present for treatment of diseases caused by expression of the bcl-2 gene
Patent: US 5750390-A 20 12-MAY-1998;
Location/Qualifiers
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ilarity 100.0%; Pred. No. 1.7e+04;
Conservative 0; Manatches 0;
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Best Local Similarity 100.0%; Pred. Mo. 1.7e+04;
Matches 35; Conservative 0; Mixmatches 0;
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Sequence 20 from patent US 5750390.
AR007306
AR007306.1 GI:3966790
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Regulation of bcl-2 gene expression
Patent: US 5831066-A 2 03-NOV-1998;
Location/Qualifiers
1. 35
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AR052604
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/organism="unknown"
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Location/Qualifiers
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100.08; Pr
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Best Local Similarity 100.0
Matches 36; Conservative
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AR052604
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1 (bases 1 to 34)
Thompson, J. D. and Draper, K. G.
Thompson, J. D. and Lager, K. G.
Method and reagent for treatment of diseases caused by expression of the bol-2 gene
Patent: US 5750390 A 16 12-MAY(1998;)
Location/Qualifiers
                                                                                                                                                     Antisense oligonucleotides inhibiting human bc1-2 gene expression Patent: US 5734033-A 2 31-MAR-1998; [Location/Qualifiers] 1. 35
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1 (bases 1 to 34)
Thompson, J.D. and Draper, K.G.
Method and reagent for treatment of diseases caused by expression
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Pred. No. 2.6e+04;
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                         Sequence 2 from patent US 5734033.
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8 c /13 g
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196083.1 GI:3940553
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l (bases 1 to 33)
Capaccioli,S., Morelli,S. and Nicolin,A.
AN ANTISENSE TRANSCRIPT ASSOCIATED TO TUMOR CELLS HAVING A T(14;18)
AN ANTISENSE TRANSCRIPT ASSOCIATED TO TUMOR CELLS HAVING A T(14;18)
TRANSLOCATION AND OLIGODEOXYNUCLEOTIDES USEFUL IN THE DIAGNOSIS AND TREATMENT OF SAID TUMOR CELLS
PATENT: WO 9627663-A 43 12-SER-1996;
CONSIGLIO NAZIONALE RICERCHE (IT)
Other Publication AN 4944296 966973.
Location/Qualifiers
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Thompson, J.D. and Draper, K.G.
Thompson, J.D. and praper for treatment of diseases caused by expression of the bcl-2 gene
Patent: US 5750390-A 5 12-MAY-1998;
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Pred. No. 2.6e+04;
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of the bcl-2 gene
Patent: US 5750390-A 21 12-MAY-1998;
Location/Qualifiers
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Best Local Similarity 100.0%; Pted. No. 4e-
Matches 33; Conservative () Mismatches
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Sequence 43 from Patent WO9627663.
A56864
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Local Similarity 100.0%;
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0.6%; Score 33; DB 6; Length 33;

Query Match

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Unclassified.

I (bases 1 to 33)

Capacioli, S. Morelli, S. and Nicolin, A.
Antisense transcript associated to tumoy cells having a T(14;18)
translocation and oligodeoxynucleotides useful in the diagnosis and
treatment of said tumor cells
treatment of said tumor 1: 1 DEC-1999;
Patent: US 6005095-A 4 21-DEC-1999;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 36)
Shir,A. and Levitzky,A.
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Pred. No. 4.5e+04;
0; Mismatches 2; Indels
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Pred. No. 4e+04;
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Sequence 44 from patent US 6005095.
AR096260
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/organism_"Homo sapiens"
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AR096260
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Capaccioli, S., Morelli, S., and Nicolin, A.
AN ANTISENSE TRANSCRIPT ASSOCIATED TO TUMOR_CELLS HAVING A T(14:18)
TRANSLOCATION AND OLIGIDENX WICKEDTIDES UŞEFUL IN THE DIAGNOSIS AND
TREATMENT OF SAID TUMOR CELLS
PATENTENT OF SAID TUMOR CELLS
PATENTENT OF SAID TUMOR (ELLS)
CONSIGLIO NAZIONALE RICERCHE (1T)
Other publication AU 4944296 960923.
LOCALION (QUALIFIERS)
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Capacciol, S., Morelli, S. and Nicolin, A.
Antisense transcript associated to tumor cells having a T(14;18)
translocation and oligodeoxynucleotides useful in the diagnosis and
treatment of said tumor cells
Patent: US 6005095-A 19 24-DEC-1999;
Location/Qualifiers
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Sequence 19 from patent US 6005095,
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Sequence 23 from patent US 6005095.
AR096239
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A56840.1 GI:3712852
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Capaccioli, S., Morelli, S. and Nicolin, A. Antisense transcript associated to tumor cells having a T(14,18) translocation and oligodeoxynucleotides useful in the diagnosis and treatment of said tumor cells Patent: US 6005095-A 23 21-DEC-1999;
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Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same method of using the same Patent: US 5582979-A 415 10-DEC-1996;
Location/Qualifiers
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Thompson,J.D. and Draper,K.G.
Method and reagent for treatment of diseases caused by expression of the bcl.2 gene
Patent: US 5750390-A 8 12-MAY-1998;
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Matches 30; Conservative 0)
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Unclassified.

1 (bases 1 to 30)

Capaccioli,S., Morelli,S. and Nicolin,A.

Antisense transcript associated to tumor cells having a T(14;18)

translocation and oligodeoxynucleotides useful in the diagnosis and
translocation and tumor cells

transment of said tumor cells

Patent: US 6005095-A 20 21-DEC-1999; Capaccioli, S., Morelli, S. and Nicolin, A. Antisense transcript associated to tumor cells having a T(14;18) translocation and oligoeoxynucleotides useful in the diagnosis and treatment of said tumor cells Patent: US 6005095-A 24 21-DEC-1999; PAT 08-SEP-2000 PAT 08-8EF-2000 PAT 18-JUN-2001 ; 0 ö 0.6%; Score 28; DB 6; Length 30; 100.0%; Pred. No. 3.7e+05; DB 6; Length 30; 3.7e+05; linear 0; Indels PST3 short site 1460 E32209 38 bp DNA Method for isolating satellite sequence. DNA Query Match 0.6%; Score 28; DB Best Local Similarity 100.0%; Pred. No. 3.7 Matches 28; Conservative 0; Mismatches Sequence 24 from patent US 6005095.
AR096240
AR096240.1 GI:10024866 Local Similarity 100.0%; Pred. No. 3.7 nes 28; Conservative 0; Mismatches 30 bp 20 from patent US 6005095. 4504 CITCAGGGTCTTCCTGAAATGCAGTGGT 4531 4504 CTTCAGGGTCTTCCTGAAATGCAGTGGT 4531 30 CTTCAGGGTCTTCCTGAAATGCAGTGGT 3 30 CITCAGGGTCTTCCTGAAATGCAGTGGT 3 Location/Qualifiers Location/Qualifiers /organism="unknown" /organism="unknown" 7 9 Sequence 20 from patent AR096236 AR096236.1\GI:10024858 Unknown. Unclassified. 9 a AR096236 Unknown. Unknown. Query Match 10 P DEFINITION ACCESSION VERSION RESULT 22 AR096236/c RESULT 23 AR096240/c DEFINITION DEFINITION ORGANISM BASE COUNT ORIGIN ORGANISM BASE COUNT Matches JOURNAL JOURNAL ACCESSION ACCESSION VERSION KEYWORDS REFERENCE REFERENCE RESULT 24 KEYWORDS SOURCE δ 1 (bases 1 to 30)
Capacatoli,S., Morelli,S. and Nicolin,A.
AN ANTISENSE TRANSCRIPT ASSOCIATED TO TUMOR CELLS HAVING A T(14;18)
TRANSLOCATION AND OLIGODEOXYMOCLECTIDES USEFUL IN THE DIAGNOSIS AND
TREATMENT OF SAID TUMOR CELLS
PATENTENT OF SAID TUMOR CELLS
PATENTENT OF SAID TUMOR CELLS
CONSIGINO NAZIONALE RICERCHE (17)96;
CONSIGINO NAZIONALE RICERCHE (17)
Other publication AU 4944296 960923. capaccioli,S., Morelli,S. and Nicolin,A.
AN ANTISENSE TRANSCRIPT ASSOCIATED TO TUMOR CELLS HAVING A T(14;18)
TRANSLOCATION AND OLICODEDXYNUCLEOTIDES USEFUL IN THE DIAGNOSIS AND
TREATMENT OF SAID TUMOR CELLS
PATENT:WO 8627663-A 23 12-28P-1996;
CONSIGLIO NAZIONALE RICECHE (IT)
Other publication AU 4944296 960923. 03-MAR-1998 PAT 03-MAR-1998 0; Gaps Gaps . 0 Score/28; DB 6; Length 28; Pred. No. 3.7e+05; Mismatches 0; Indels linear Length 30; Length 30; 0; Indels Query Match 0.6%; Score 28; DB 6; Lk Best Local Similarity 100.0%; Pred. No. 3.7e+05; Matches 28; Conservative 0; Mismatches 0; Query Match 0.6%; Score 28; DB 6; Ls Best Local Similarity 100.0%; Pred. No. 3.7e+05; Matches 28; Conservative 0; Mismatches 0; 30 bp Seguence 23 from Patent W09627663. A56844 A56844 GI:3712856 A56841 30 bp Sequence 20) from Patent W09627663 /organism="unidentified" /db_xref="taxon:32644" 9 c 7 g /organism="unidentified" /db_xref="taxon:32644" 9 c 7 g E 4504 CTTCAGGGTCTTCCTGAAATGCAGTGGT 4531 1 CCTGTGGTCCACCTGGCCCTCGGCCAAG 28 Location/Qualifiers 30 CTTCAGGGTCTTCCTGAAATGCAGTGGT 3 Location/Qualifiers 100.0%; Pi GI:3712853 Query Match 0.0%; Best Local Similarity 100.0% Matches 28; Conservative 1729 CCTGTGGTCCACCTGGCCCTG (bases 1 to 30) unidentified unclassified. unidentified unclassified. unidentified unidentified ORGANISM ORGANISM DEFINITION BASE COUNT ACCESSION VERSION BASE COUNT ORIGIN RESULT 20 A56841/c DEFINITION ACCESSION FERENCE AUTHORS JOURNAL RESULT 21 REFERENCE AUTHORS JOURNAL KEYWORDS A56844/c KEYWORDS FEATURES FEATURES COMMENT COMMENT ORIGIN g

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                                       Eukaryota; Metazoa; Mallusca; Gastropoda; Archaeogastropoda;
Halotidae; Hallotis.

E 1 (bases 1 to 38)

E 1 (bases 1 to 38)

E Hideaki, T. and Masabi, S.

Method for isolating satellite sequence

L Patent: JP 2000060559-A 11 29-FEB-2000;

NATL INST OF AGROBIOLOGICAL RESOUCES
OS Hallotis discus show JP 2000060559-A/11

PD 29-FEB-2000

PF 18-AUG-1998 JP 1998232153

PR HIDEAKI TAKAHASHI, MASASHI SEKINO

PC C12N15/09,C12Q1/68,C12N15/00

CC FF Source 1. 38

Location/Qualifiers

FT Source //organism='Hallotis discus discus'.
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Pred. No. 5.5e+05;
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/organism="Haliotis discus discus"
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Best Local Similarity 93.5%; Pred. No. 4.2e+05;
Matches 29; Conservative 0; Mismatches 2;
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Sequence 403 from patent US 5582979.
131491 GI:1822282
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Haliotis discus discus.
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Weber, J.L.
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Matches 32; Conservative
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1 (bases 1 to 40)
Weber, J.L.
Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 403 10-DEC-1996;
Location/Qualifiers
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1 (bases 1 to 27)
Thompson, J.D. and Draper, K.G.
Method and reagent for treatment of diseases caused by expression of the bci-2 gene
Patent: US 5750390-A 17 12-MAY-1998;
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Pred. No. 5.5e+05;
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PAT 18-JUN-2001

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PAT 06-FEB-1997

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1 (bases 1 to 31)
Weber, J.L.
Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 271 10-DEC-1996;
Location/Qualifiers
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JP 2000060559-A/5.
Hallotis discus discus.
SM Hallotis discus discus.
Eukaryota; Metazca; Mollusca; Gastropoda; Archaeogastropoda;
Eukaryota; Metazca; Mollusca; Gastropoda; Archaeogastropoda;
Hallotidae; Hallotis.
E 1 (bases 1 to 32)
S Hideaki,T. and Masashi,S.
I (bases 1 to 32)
S Hideaki,T. and Masashi,S.
L Patent: JP 2000060559-A 5 29-FEB-2000;
NATL INST OF AGROBIOLOGICAL RESOUCES
OF Hallotis discus discus
PN JP 2000060559-A/5
PD 29-FEB-2000
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/organism='Haliotis discus discus'.
Location/Qualifiers
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/organism="Haliotis discus discus"
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Pred. No. 6.4e+05;
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Sequence 210 from patent US 5582979.
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Best Local Similarity 93.3%;
Matches 28; Conservative 0
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1 (base; 1 to 31)
Weber,J.L.
Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same Patent: US 5582979-A 215 10-DEC-1996;
Location/Qualifiers
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Sequence 271 from patent US 5582979.
131359.1 GI:1822150
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Best Local Similarity 93.3%; Pred. No. 6.4e+05;
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Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 376 10-DEC-1996;
Location/Qualifiers
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                                  1 (bases 1 to 32)
Weber, J.L.
Length polymorphisms in (dc-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 55812979-A 210 10-DEC-1996;
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I31147.1 GI:1821938
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Weber, J.L.
Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same method of using the same patent: US 5582979-A 110 10-DEC-1996;
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Weber, J.L.
Ength polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 218 10-DEC-1996;
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                                                                                                                            Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 172 10-DEC-1996;
Location/Qualifiers
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Weber,J.L.
Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 180 10-DEC-1996;
Location/Qualifiers
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Patent: US 5582979-A 192 10-DEC-1996;
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Best Local Similarity 93.3%;
Matches 28; Conservative
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Weber, J.L.
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Weber, J.L.
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Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 322 10-DEC-1996;
Location/Qualifiers
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Weber,J.L.
Length bolymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 77 10-DEC-1996;
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Sequence 322 from patent US 5582979.
131410 GI:1822201
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Location/Qualifiers
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Weber,J.L.
Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 104 10-DEC-1996;
Location/Qualifiers
                                                                                                                                    Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same Patent: US 5582979-A 189 10-DEC-1996;
Location/Qualifiers
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93.3%; Pred. No. 6.5e+05;
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Sequence 358 from patent US 5582979.
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131446.1 GI:1822237
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Sequence 189 from patent US 5582979.
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Sequence 104 from patent US 5582979.
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Best Local Similarity 93.39
Matches 28, Conservative
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1 (bases 1 to 35)
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Weber, J.L.
Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same method of using the same Patent: US 55812979-A 140 10-DEC-1996;
Location/Qualifiers
Weber, J.L.
Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and
method of using the same
method of using the same
Patent: US 5582979-A 358 10-DEC-1996;
Location/Qualifiers
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Best Local Similarity 93.3%; Pred. No. 6.5e+05;
Matches 28; Conservative 0; Mismatches 2; Indels
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/db_xref="taxon:9823"
/note="Microsatellite"
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Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and
method of using the same
Patent: US 5582979-A 68 10-DEC-1996;
Location/Qualifiers
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Weber, J.L.
Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 198 10-DEC-1996;
Location/Qualifiers
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Sequence 68 from patent US 5582979.
I31156.1 GI:1821947
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Weber, J. L.
Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 397 10-DEC-1996;
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O.5%; Score 26.8; DB 6;
Best Local Similarity 93.3%; Pred. No. 6.5e+05;
Matches 28; Conservative 0; Mismatches 2;
Sequence 397 from patent US 5582979.
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26-MAY-1994 (first entry)

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B-cell acute lymphocytic leukemia; breast cancer; colon carcinoma; neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif; hairpin; hepatitis delta virus; group I intron; RNaseP; ss.
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Homo sapiens.

W09323057-A

25-NOV-1993

93WO-US04573 13-MAY-1993;

14-MAY-1992 26-AUG-1992

92US-0936110. 92US-0936421. 92US-0936422. 92US-0936531. 92US-0936532. 93US-0008910 93US-0006122 26-AUG-1992; 26-AUG-1992; 26-AUG-1992; 26-AUG-1992; 07-DEC-1992; 19-JAN-1993;

(RIBO-) RIBOZYME PHARM INC.

Draper KG, Thompson JD;

WPI; 1993-386203/48

New enzymatic RNA molecules (ribozymes) - which cleave mRNA associated with tumours or mRNA expressed from gene encoding multiple drug resistance

Claim 3; Fig 6; 69pp; English.

Sassociated with development or maintenance of Chronic myelogenous lettermia (CML), promyelocytic leukemia, Burkitt's lymphoma, or acute lymphocytic leukemia, follicular lymphoma, B-cell acute well and procytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, breast cancer, colon carcinoma neuroblastoma and lung cancer. The full length mRNAs containing these target sequences, encode aberant cellular proteins which are able to control cellular proliferation and are directly linked to a leukemic phenotype. These target sequences are identified by the ribozyme of the invention. The ribozymes is formed in a hammerhead motifi, but may also be formed in the motifi of a hairpin, hepatitis delta virus, group I intron or RNaseP-like RNA. These ribozymes may be used to inhibit the development or expression of a transformed phenotype in man and other animals by modulating expression of the corresponding gene. Cleavage of target mRNAs expressed in pre-neoplastic and transformed cells alicitize inhibition of the transformed state. Multiple drug resistance (mdr-1) mRNA specific ribozymes remove the mechanism of the ranspales for tumours. The ribozymes remove the mechanism of the ranspales for tumours. The ribozymes may also be used to study sequences given in AAQ51825-2266 represent areas of mRNAs which are genetic drift and mutations within cells. 

Sequence 37 BP; 2 A; 12 C; 12 G; 11 U; 0 other;

Gaps ; Length 37; Indels Ouery Match 0.7%; Score 37; DB 14; Best Local Similarity 70.3%; Pred. No. 1.2e+02; 11; Mismatches 26; Conservative Query Match Matches

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1415 GGCGAGAGGTGCCGTTGGCCCCGGTTGCTTTTCCTCT 1451 1 GGCGAGGGCCCUUGCCCCUUGCCUCU 37 δ g

AAQ51961 standard; RNA; 36 BP AAQ51961; RESULT 2 **AA**Q51961

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resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin D; vinblastine; small intestine; kidney; adrenal gland; adenocarcinoma; bowel; transformed phenotype; promyelocytic leukemia; human; chronic myelogenous leukemia; CML; follicular lymphoma; B-cell acute lymphocytic leukemia; breast cancer; colon carcinoma; neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif; hairpin; hepatitis delta virus; group I intron; RNasep; ss.
                                  Multiple drug resistance; mdr-1; ribozyme; membrane protein; liver;
                BCL-2 mRNA ribozyme cleavable nucleotide (1997).
                                                                                                                                                                                                   920S-0936110.
920S-0936421.
920S-0936422.
920S-0936531.
920S-0936532.
920S-098131.
                                                                                                                                                                                                                                                                  93US-0008910.
                                                                                                                Homo sapiens.
                                                                                                                                                                  13-MAY-1993;
                                                                                                                                WO9323057-A.
                                                                                                                                                                                                                                                                  19-JAN-1993;
                                                                                                                                                 25-NOV-1993.
                                                                                                                                                                                                     26-AUG-1992,
                                                                                                                                                                                             -MAY-1992
                                                                                                                                                                                                                       26-AUG-1992
                                                                                                                                                                                                                               26-AUG-1992
                                                                                                                                                                                                                                         26-AUG-1992
                                                                                                                                                                                                                                                07-DEC-1992
                                                                                                                                                                                                                                                         9-JAN-1993
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New enzymatic RNA molecules (ribozymes) - which cleave mRNA associated with tumours or mRNA expressed from gene encoding multiple drug resistance

(RIBO-) RIBOZYME PHARM INC.

Draper KG, Thompson JD; WPI; 1993-386203/48. Claim 3; Fig 6; 69pp; English.

The sequences given in AAQ51825-2266 represent areas of mRNAs which are associated with development or maintenance of chronic myelogenous. associated with development or maintenance of chronic myelogenous. Cleukemia (CML), promyelocytic leukemia, Burkitt's lymphon, or ocute lymphocytic leukemia, follicular lymphona, B-cell acute. Iymphocytic leukemia, brast cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAs containing these target sequences, encode aberant cellular proteins which are able to control cellular proliferation and are directly linked to a leukemic phenotype. These target sequences are identified by the ribozyme of the invention. The ribozymes is formed in a hammerhead motif, but may also be formed in the motif of a hairph, hepatitis delta virus, group intron or RNaseP-like RNA. These ribozymes may be used to inhibit the development or expression of a transformed phenotype in man and other animals by modulating expression of the corresponding gene. Cleavage of target mRNAs expressed in pre-neoplastic and transformed cells alicits inhibition of the transformed state. Multiple drug resistance (mdr-1) mRNA specific ribozymes remove the mechanism of the ribozymes may also be used to study genetic drift and mutations within cells.

Sequence 36 BP; 10 A; 13 C; 8 G; 5 U; 0 other;

Gaps .; 0 Length 36; Indels // Match 0.7%; Score 36; DB 14; L
Local Similarity 86.1%; Pred. No. 1.9e+02;
nes 31; Conservative 5; Mismatches 0; Matches

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1997 ACCTGAACCGCACCTGCACACCTGGATCCAGGATA 2032 ŏ

1 ACCUGAACCGGCACCUGCACCUGGAUCCAGGAUA 36

셤

AAX21301 standard; RNA; 36

RESULT 3 AAX21301 AAX21301;

1 14:55:53 2003

Sun Jun

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/bound_molety= "bases 11-17 of AAX21302"
                                                                  32..36
/*tag= c
/bound_moiety= "bases 3-7 of AAX21302"
                                                                                                                                                                                                                                                                       1 GAAAAACCUGGAUGUCACUGGCCACUGAGGAGGUUU 36
                                                                                                         98WO-US12570.
                                                                                                                   97US-0879078
                                                             /*tag= b
                                                                                                                             (INNO-) INNOVIR LAB INC
                                                                                                                                                                                                                            the function of a gene.
                                                                                                                                               WPI; 1999-095336/08.
                                                                                                                   19-JUN-1997;
                                                                                                         16-JUN-1998;
                                                                   misc_binding
                                                                                     WO9858057-A1
                                                                                               23-DEC-1998.
                                                                                                                                      Ludwig J,
                                                         stem_loop
                                                                                                                                                                                                                                                                                          RESULT 4
AAQ51966
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The invention relates to a composition that induces cleavage of an RNA substrate by forming a structure similar to a hammerhead ribozyme. This sequence represents a fragment of the human bcl-2 mRNA which binds the catalytic RNA AAX21302. Compositions containing catalytic RNAs can be used in pharmaceutical compositions to specifically inactivate the expression of genes in eukaryotes, prokaryotes and viruses, especially of human genes, e.g. tumour genes or viral genes or RNA molecules in a cell. They can also be used for inactivation of the expression of plant or insect genes, and used as pesticides. They can be used to identify
Compositions inducing cleavage of an RNA substrate - useful as drugs or pesticides, and for identifying the function of a gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 36 BP; 10 A; 8 C; 10 G; 8 U; 0 other;
                                                                                                                                                                                            Example 3; Fig 5A; 62pp; English.
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Gaps .; 0 Length 36; Indels ; 0 Query Match

0.7%; Score 36; DB 20;
Best Local Similarity 77.8%; Pred. No. 1.9e+02;
Matches 28; Conservative 8; Mismatches 0;

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4147 GAAAAACCTGGATGTCACTGGCCACTGAGGAGCTTT 4182

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AAQ51966 standard; RNA; 35 BP
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Multiple drug resistance; mdr-1; ribozyme; membrane protein; liver; resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin p; vinblastine; small intestine; kidney; adrenal gland; adenocarcinoma; bowel; transformed phenotype; promyelocytic leukemia; human; chronic myelogenous leukemia; CML; follicular lymphocytic leukemia; becell acute lymphocytic leukemia; reasst cancer; colon carcinoma; neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif; hairpin; hepatitis delta virus; group I intron; RNaseP; ss.
                                  BCL-2 mRNA ribozyme cleavable nucleotide (2180).
                                                                                                                                                                                                                                                    920S-0936422.
920S-0936531.
920S-0936532.
920S-0987131.
930S-0006122.
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92US-0882885.
92US-0936110.
92US-0936421.
                                                                                                                                                                                                93WO-US04573
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                                                                                                                                                                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC
                (first entry)
                                                                                                                                                                                                                                                                                                                                            Draper .KG, Thompson JD;
                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                13-MAY-1993;
                  26-MAY-1994
                                                                                                                                                            WO9323057-A.
                                                                                                                                                                                                                                                                                                       19-JAN-1993;
                                                                                                                                                                              25-NOV-1993
                                                                                                                                                                                                                             14-MAY-1992
                                                                                                                                                                                                                                                                                     07-DEC-1992
                                                                                                                                                                                                                                     26-AUG-1992
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                                                                                                                                                                                                                                                          26-AUG-1992
                                                                                                                                                                                                                                                                  26-AUG-1992
                                                                                                                                                                                                                                                                           26-AUG-1992
aaq51966;
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Cleavage; hammerhead ribozyme; catalytic RNA; gene expression; ss; eukaryote; prokaryote; virus; human; tumour; plant; insect; pesticide;

Location/Qualifiers

Homo sapiens

/*tag=

misc_binding

Human bcl-2 mRNA as catalytic RNA substrate.

11-MAY-1999 (first entry)

The sequences given in anylosis-zero interest areas of anthon and associated with development or maintenance of chronic myelogenous associated with development or maintenance of chronic myelogenous leakemia (CML), promyelocytic leukemia, follicular lymphoma, B-cell acute caute lymphocytic leukemia, follicular lymphoma, B-cell acute acute lymphocytic leukemia, follicular lymphoma, B-cell acute acute lymphocytic leukemia, breast cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAs containing these target sequences, encode aberant cellular proteins which are able to control sequences. These target sequences are identified by the ribozyme of the invention. The motif of a hairpin, hepatitis delta virus, group also be formed in the motif of a hairpin, hepatitis delta virus, group in intron or RNAsseP-like RNA. These ribozymes may be used to inhibit the development or expression of a transformed phenotype in man and other animals by modulating expression of the corresponding gene. Cleavage of target mRNAs expressed in pre-neoplastic and transformed cells elicits inhibition of the transformed state. Multiple drug resistance used by transformed cells and thus enhances drug therapies for tumours. The ribozymes remove the mechanism of therapies for tumours. The ribozymes may also be used to study therefore drift and mutations within cells. The sequences given in AAQ51825-2266 represent areas of mRNAs which are Claim 3; Fig 6; 69pp; English. multiple drug resistance

associated with tumours or mRNA expressed from gene encoding New enzymatic RNA molecules (ribozymes) - which cleave mRNA

WPI; 1993-386203/48.

Gaps ö Query Match

0.7%; Score 35; DB 14; Length 35;
Best Local Similarity 85.7%; Pred. No. 3.2e+02;
Matches 30; Conservative 5; Mismatches 0; Indels Sequence 35 BP; 14 A; 10 C; 6 G; 5 U; 0 other;

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AAQ8664

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This is a human bc1-2 oligonucleotide based on which an antisense oligonucleotide complementary to the translation initiation site of the human bc1-2 mRNA can be constructed. Bc1-2 antisense oligonucleotides strategic sites such as the translation initiation site, donor and acceptor splicing sites, or sites for transportation or degradation. Blocking translation at such strategic sites prevents the formation of a functional bc1-2 gene product. These oligonucleotides may be used for treating cancers associated with high levels of bc1-2 gene expression, especially lymphomas and some leukaemias.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B cell lymphoma/leukaemia-2; bcl-2; oncogene; antisense; lymphoma; leukaemia; colon carcinoma; rectal carcinoma; pancreatic cancer; breast cancer; ovarian cancer; prostate cancer; renal cell carcinoma; hepatoma; bile duct carcinoma; choriocarcinoma; cervical cancer; testicular cancer; lung carcinoma; bladder carcinoma; melanoma; head and neck cancer; brain cancer; cytostatic; human; gene therapy;
                                                                                                                                                                                                                                                                                                                     Anti-sense oligo:nucleotide(s) complementary to BCL-2 mRNA - useful for treating cancers, e.g. lymphoma(s) and some leukaemia(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C; 13 G; C.
0.7%; Score 35; DB 19; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 35 BP; 6 A; 8 C; 13 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1442 CTTTTCCTCTGGGAAGGATGGCGCACGCTGGGAGA 1476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CTTTTCCTCTGGGAAGGATGGCGCACGCTGGGAGA 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0%; Pred. No. 3.2
Matches 35; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Columns 3-4; 21pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bcl-2 antisense oligonucleotide.
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2000US-237009P.
2000US-0709170.
                                                                                                 94US-0288692.
                                                                                                                                                    88US-0288692.
94US-0217082.
                                                                                                                                     92US-0840716.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABL54150 standard; DNA; 35
                                                                                                                                                                                                          (UYPE-) UNIV PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                 WPI; 1998-229881/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200217852-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-AUG-2000; 29-SEP-2000; 10-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                 24-MAR-1994;
                                                                                                                                                    22-DEC-1988;
24-MAR-1994;
                                                                                                                                     21-FEB-1992;
                        US5734033-A.
                                                               31-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-MAR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABL54150;
                                                                                                                                                                                                                                              Reed J:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 7
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                                                                                                                                                                                                                                                                         Anticode oligomer; antisense oligonucleotide; bcl-2; cancer; therapy; leukemia; lymphoma; solid tumor; breast cancer; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The antisense oligonucleotide TI-AS (AA086643) straddles the translation-initiation site in the mRNA coding strand of the human bcl-2 gene and is complementary to this region. It reduces the expression of bcl-2 gene product thereby inducing programmed cell death of certain cancer cells. The corresp. sense bcl-1 sequence was synthesized for use as a control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense oligonucleotide; bcl-2 gene; lymphoma; leukaemia; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anti-code oligomers which bind to bcl-2 mRNA - for the treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 35; DB 16; Length 35; 100.0%; Pred. No. 3.2e+02;
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2180 GTCAACATGCCTGCCCCAAACAAATATGCAAAAGG 2214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 35 BP; 6 A; 8 C; 13 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1442 CTTTTCCTCTGGGAAGGATGGCGCACGCTGGGAGA 1476
                     1 CTTTCCTCTGGGAAGGATGCCGCACGCTGGGAGA 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Pred. No. 3.2 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of human solid tumours, esp. breast cancer
                                                                                                                                                                                                                                   Bcl-2 translation initiation region.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             94WO-US10725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93US-0124256
                                                                                                                            AAQ86644 standard; DNA; 35
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                                                                                                                                                                                                  27-SEP-1995 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1995-139394/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
les 35; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (REED/) REED J C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                    WO9508350-A
                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAR-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer; ss.
                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
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Matches

δ g RESULT 6

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Gaps

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Length 35; Indels us-09-375-514-19.rng

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                                                                                                                    Treating or preventing cancer, tumors and carcinomas, comprises administering B cell lymphoma/leukemia-2 antisense oligonucleotide at high doses for short period for time with one or more cancer
                                                                                                                                                                                                                                                                  The present sequence is that of a B cell lymphoma/leukaemia-2 (bcl-2) antisense oilgonucleotide. The present invention is directed to the use of bcl-2 antisense oilgomers, particularly G3139 (see ABL54148), to treat and prevent bcl-2 related disorders. Administration at high doses results in significant therapeutic responses, including low toxicity, high tolerance and prolonged survival. Administration at high doses for short periods of time (less than 14 days) also provides significant therapeutic responses in the treatment of cancer. The bcl-2 antisense oligomer may also be used to increase the sensitivity of a subject to cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      reproductive system, genitourinary system, breast, endocrine system, brain, central nervous system, peripheral nervous system, kidney, lung, respiratory system, thorax, gastrointestinal and all mentary canal, lymph nodes, pancreas, hepatobiliary system, or cancer of unknown primary site, non-Hodgkin's lymphoma, Hodgkin's lymphoma, leukaemia, colon carcinoma, rectal carcinoma, pancreatic, brain cancer, renal cell carcinoma, hepatoma, bile duct carcinoma, choricoarcinoma, ung carcinoma, hepatoma, bile duct carcinoma, choricoarcinoma, lung carcinoma, bladder carcinoma and melanoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               therapeutics, and in combination with hormone treatment or gene therapy. Conditions that may be treated or prevented include cancer of the haematopoietic system, skin, bone and soft tissue,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 35;
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; Pred. No. 3.2e+02;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BCL-2 mRNA ribozyme cleavable nucleotide (2043).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1442 CTTTTCCTCTGGGAAGGATGGCGCACGCTGGGAGA 1476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 35 BP; 6 A; 8 C; 13 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 CTTTTCCTCTGGGAAGGATGGCGCACGCTGGGAGA 35
                                                                                                                                                                                                                              Disclosure; Page 53; 64pp; English.
                                       Fingert H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ51962 standard; RNA; 34 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 100.0 Matches 35; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAY-1994 (first entry)
                                       Warrel RP, Klem RE,
                                                                             WPI; 2002-371796/40.
(GENT-) GENTA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        all claimed).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                       therapeutics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ51962;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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The sequences given in AAQ51825-2266 represent areas of mRNAs which are associated with development or maintenance of chronic myelogenous associated with development or maintenance of chronic myelogenous leukemia, promyelocytic leukemia, follicular lymphoma, B-cell acute cacute lymphocytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, breast cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAs containing these target sequences, encode aberant cellular proteins which are able to control cellular proliferation and are directly linked to a leukemic phenotype. These target sequences are identified by the ribozyme of the invention. The ribozymes is formed in a hammerhead motif, but may also be formed in the motif of hairph, hepatitis delta virus, group intron or RNaseb-like RNA. These ribozymes may be used to inhibit the development or expression of a transformed phenotype in man and chear animals by modulating expression of the corresponding gene. Cleavage of target mRNAs expressed in pre-neoplastic and transformed cells elicits inhibition of the transformed state. Multiple drug frug resistance used by transformed cells and thus enhances drug therapies for tumours. The ribozymes may also be used to study
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin D; vinblastine; small intestine; kidney; adrenal gland; adenocarcinoma; bowel; transformed phenotype; promyelocytic leukemia; human; chronic myelogenous leukemia; CML; follicular lymphoma; B-cell acute lymphocytic leukemia; breast cancer; colon carcinoma;
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                                                                                                                                                                                                                                                                                                                                                                  New enzymatic RNA molecules (ribozymes) - which cleave mRNA associated with tumours or mRNA expressed from gene encoding
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Pred. No. 5.2e+02;
8; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3; Fig 6; 69pp; English.
                                                                                                                                 920S-0936531.
920S-0936532.
920S-0987131.
930S-0006122.
930S-0008910.
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Local Similarity 76.5%;
les 26; Conservative (
                                    92US-0882885.
92US-0882885.
92US-0936110.
  93WO-US04573.
                                                                                              92US-0936421
92US-0936422
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                                                                                                                                                                                                                                                                                                                                                                                                                multiple drug resistance
                                                                                                                                                                                                                                                                                               Draper KG, Thompson JD;
                                                                                                                                                                                                                                                                                                                                    WPI; 1993-386203/48.
13-MAY-1993;
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19-JAN-1993;
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                                                                             26-AUG-1992;
                                                                                                                                       26-AUG-1992;
                                      14-MAY-1992
14-MAY-1992
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neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif; hairpin; hepatitis delta virus; group I intron; RNaseP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          other animals by modulating expression of the corresponding gene. Cleavage of target mRNAs expressed in pre-neoplastic and transformed cells elicits inhibition of the transformed state. Multiple drug resistance (mdr-1) mRNA specific ribozymes remove the mechanism of
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73.5%; Pred. No. 5.2e+02;
tive 9; Mismatches 0; Indels
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920S-0936110.
920S-0936421.
920S-0936422.
920S-0936531.
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ches 25; Conservative
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                                                                                                     Homo sapiens
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19-JAN-1993;
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26-AUG-1992
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The sequences given in AAQ51825-2266 represent areas of mRNAs which are associated with development or maintenance of chronic myelogenous leukemia (CML), promyelocytic leukemia, Burkitt's lymphona, or cute lymphocytic leukemia, follicular lymphona, B-cell acute lymphocytic leukemia, follicular lymphona, B-cell acute lymphocytic leukemia, beast cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAs containing these target sequences, encode aberant cellular proteins which are able to control cellular proliferation and are directly linked to a leukemic phenotype. These target sequences are identified by the ribozyme of the invention. The ribozymes is formed in a hammerhead motif, but may also be formed in the motif of a halitpin, hepatitis delta virus, group intron or RNaseP-like RNA. These ribozymes may be used to inhibit the development or expression of a transformed phenotype in man and other animals by modulating expression of the corresponding gene. Cleavage of target mRNAs expressed in pre-neoplastic and transformed cells elicits inhibition of the transformed state. Multiple drug frug resistance used by transformed cells and thus enhances drug theraples for tumours. The ribozymes may also be used to study
                                               Multiple drug resistance; mdr-1; ribozyme; membrane protein; liver; resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin D; vinblastine; small intestine; kidney; adrenal gland; adenocarcinoma; bowel; transformed phenotype; promyelocytic leukemia; human; chronic myelogenous leukemia; CML; follicular lymphoma: becall acute lymphocytic leukemia; brast cancer; colon carcinoma; neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif; hairpin; hepatitis delta virus; group I intron; RNaseP; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New enzymatic RNA molecules (ribozymes) - which cleave mRNA associated with tumours or mRNA expressed from gene encoding multiple drug resistance
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               BCL-2 mRNA ribozyme cleavable nucleotide (1592).
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Pred. No. 8.4e+02;
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92US-0936422.
92US-0936531.
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92US-0987131.
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84.8%;
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les 28; Conserv
                                                                                                                                                                                                                            Homo sapiens.
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RESULT 12

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                                                                                                                                                                                                                                                                                                                           genetic locus;
                                                                                                                                                                                                                                                                                                                                                  antisense; therapy; proliferative disorder; neoplastic disease; psortasis; vasculogenesis; anglogenesis; apoptosis; cytostatic; bcl-2; immunoglobulin heavy chain; IgH; fusion protein; DCLL; follicular lymphoma; FT; Chromosomal translocation; ds.
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                                                                                                                                                                                                                                                                                                                           double stranded RNA dependent protein kinase; PKR;
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20..21
/*tag- a
/note- "Translocation occurring site"
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14.4%; Pred. No. 9.7e+02;
ve 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4098 CAAAAGCATTCTGAGAAGGTGAGATAAGCCCTGAGT 4133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 12; 54pp; English.
                                                             AAD12894 standard; DNA; 36 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-JAN-2001; 2001WO-IL00094
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-JAN-2000; 2000US-0179361.
22-DEC-2000; 2000US-0258010
                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity 94.4
nes 34; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shir A, Levitzky A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-488878/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200157205-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                          16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        population
                                                                                                                          AAD12894;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                           Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
RESULT 11
                                 AAD12894
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1 CAAAAGCATTCTGAGAAGGTGAGATAAGAACTGAGT 36

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a method for selective killing of cells in a targetted cell population by selectively activating double stranded (ds) RNA dependent protein kinase (PRR). The method involves selecting sequence at single genetic locus in targetted cell population that is absent from equivalent locus in non-targetted cell population, obtaining anti-sense RNA having sequence homology with the genetic locus, and permitting anti-sense RNA to hybridise with the RNA transcribed from the permitting anti-sense RNA to hybridise with the RNA transcribed from the slow used for treating proliferative discorders such as neoplastic disease, psoriasis and vasculogenesis or andiogenesis. The present sequence is an anti-sense RNA targetted to antiapoptotic bloll-limmonglobulin heavy chain (IGH) locus fusion gene. The DCLL and follicular lymphoma (FT) is characterised by a t(14;18)(q32;q21) chromosomal translocation of bcl-2-IGH fusion gene.
                                                                                                                                                      Human, double stranded RNA dependent protein kinase; PKR; genetic locus; antisense; therapy; proliferative disorder; neoplastic disease; psortasis; vasculogenesis; anglogenesis; apoptoasis; cytostatic; bcl-2; immunoglobulin heavy chain; 199; fusion protein; DCLi; follicular lymphoma; FT; chromosomal translocation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Activating double stranded RNA dependent protein kinase in targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cell population, by hybridizing antisense RNA with sequence at s
genetic locus in the population, that is absent in non-targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                        Anti-sense RNA targetted to bcl-2-IgH fusion gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.6%; Score 31.2; DB 22;
91.7%; Pred. No. 2.2e+03;
iive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (YISS ) YISSUM RES DEV CO HEBREW UNIV JERUSALEM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4098 CAAAAGCATTCTGAGAAGGTGAGATAAGCCCTGAGT 4133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 36 BP; 7 A; 11 C; 3 G; 15 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 30; Page 12; 54pp; English.
                AAD12893 standard; RNA; 36 BP
                                                                                                                                                                                                                                                                                                                                                                          31-JAN-2001; 2001WO-IL00094.
                                                                                                                                                                                                                                                                                                                                                                                                          2000US-0179361
2000US-0258010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT35442 standard; DNA; 30
                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          33; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Levitzky A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-488878/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                  WO200157205-A1
                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                            31-JAN-2000;
22-DEC-2000;
                                                                                   16-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27 - MAY - 1997
                                                                                                                                                                                                                                                                                                                                       09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          population
                                                    AAD12893;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAT35442;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shir A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 13
AAD12893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAT35442
                                   δλ
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AAT35441-T35443 are PCR primers used to amplify the chimeric bcl-2/IgH (immunoglobulin heavy chain) gene present in the t(14; 18) chromosomally translocated non-Hodgkin's lymphoma cell line DOHH-2. The transcript produced is used for the design of antisense transcripts which can inhibit the production of bcl-2 protein in t(14; 18) translocated cells and induce cell death. The antisense transcripts can be used for the monitoring treatment of t(14; 18)-associated tumours and for translocation.
                                                  chromosome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymorphism; repeat sequence; genetic marker; primer; amplification; PCR; polymerase chain reaction; paternity; maternity; human; pedigree; linkage analysis; genetic disease; animal; plant; breeding; locus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                           Non-Hodgkin's lymphoma; cell line DOHH2; DHL-4; leukaemia; chromosom t14; t18; q32; q21; N region; antisense; expression; inhibitor; bcl-2; 1gH; immunoglobulin heavy chain; hybrid; cell death; tumour; cancer; neoplasia; diagnosis; translocation; chimera; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                     New chimeric bcl-2/IgH antisense transcript of t(14; 18) translocated cells - used to develop oligodeoxynucleotides for the diagnosis and treatment of tumours associated with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
;
                 Non-Hodgkin's B cell line bcl-2/IgH chimeric gene PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.6%; Score 30; DB 17; Length 30; 100.0%; Pred. No. 3.7e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (dC-dA)n.(dG-dT)n polymorphic repeat sequence #7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 30 BP; 3 A; 9 C; 6 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Preα. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4256 GACCTIGTTTCTTGAAGGTTTCCTCGTCCC 4285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GACCTTGTTCTTGAAGGTTTCCTCGTCCC 30
                                                                                                                                                                                                                                                                                                                               Morelli S, Nicolin A;
                                                                                                                                                                                                                                                                                              (CNDR ) CONSIGLIO NAZ DELLE RICERCHE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page 11; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hybridisation; chromosome; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT66066 standard; DNA; 30 BP.
                                                                                                                                                                                                                                     96WO-EP00852.
                                                                                                                                                                                                                                                                   95IT-0MI0420.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-JUN-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 100.0
                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-425424/42.
                                                                                                                                                                                                                                                                                                                             Capaccioli S,
                                                                                                                                                                                                                                                                                                                                                                                                                                         translocation.
                                                                                                                                                                                                                                  02-MAR-1996;
                                                                                                                                                                                                                                                                   03-MAR-1995;
                                                                                                                                                                       WO9627663-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                     12-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-DEC-1996
                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT66066;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 14
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having the sequence (dC-dA)n.(dG-dT)n which can be used as genetic
markers. Frimers based on these sequences can be used to detect these
repeats, especially for use in e.g paternity or maternity testing,
human genetic analysis such as linkage analysis of genetic disease,
commercial animal or plant breeding or pedigree analysis.
The repeats, when analysed, fall into 4 categories:
The repeats which are alternating tandem CA repeats with no
interruptions and without adjacent repeats of another sequence;
The repeats separated by no more than 3 consecutive non-repeat bases;
The repeats separated by no more than 3 consecutive non-repeat bases;
The repeats separated by no more than 3 consecutive non-repeat bases;
The repeats separated by no more than 3 consecutive non-repeat bases;
This no more than 3 consecutive non-repeat bases from a run of at least
The repeats which are defined as 5 or more chan consecutive on repeat bases from a run of at least
Thin interrupted dinucleotide or longer repeats of a sequence other than
The repeats which are defined as for the perfect
CG-dA)n, CG-d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yeast Artificial Chromosome; YAC; polymerase chain reaction; PCR; sequence tagged site; genetic disorder; diagnosis; abnormality; Prader-Willi; Angelman; Beckwith-Wiedermann; syndrome; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ô
                                                                                                                                                                                                                                                                                                     Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n - using novel nucleic acid mols. as primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.6%; Score 30; DB 18; I
100.0%; Pred. No. 3.7e+03;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 30 BP; 15 A; 13 C; 2 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                             Example 8; Column 57-58; 186pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 CACACACAGACAGACACACACACACA 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers 5..38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human chromosomal repeat element.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /rpt_type= tandem
5..6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                         89US-0341562.
94US-0222177.
                   89US-0341562
                                                                   91US-0754351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ68850 standard; DNA; 38
                                                                                                                                                              (MARS-) MARSHFIELD CLINIC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100. hes 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag= b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag=
                                                                                                                                                                                                                                                        WPI; 1997-042299/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CA) 5G(ACA) G(AC) 7A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          repeat_region
                 21-APR-1989;
                                                                05-SEP-1991;
                                                                                                               04-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                         -APR-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  repeat_unit
                                                                                                                                                                                                           Weber JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ68850;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ68850
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WO9406936-A.

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The sequences given in AAQ51825-2266 represent areas of mRNAs which are associated with development or maintenance of chronic myelogenous leukemia (CMI), promyelocytic leukemia, Burkitt's lymphoma, or cute lymphocytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, follicular lymphoma, B-cell acute lymphocytic leukemia, breast cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAs. containing these target sequences, encode aberant cellular proteins which are able to control cellular proliferation and are directly linked to a leukemic phenotype. These target sequences are identified by the ribozyme of the invention. The ribozymes is formed in a hammerhead motif, but may also be formed in the motif of a hairpin, hepatitis delta virus, group in thron or RNaser-like RNA. These ribozymes may be used to inhibit the development or expression of a transformed phenotype in man and other animals by modulating expressed in pre-neoplastic and transformed cells and thus enhances drug resistance (mdr-1) mRNA specific ribozymes remove the mechanism of the resistance (mdr-1) mRNA specific ribozymes enhances drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Non-Hodgkin's lymphoma; cell line DOHH2; DHL-4; leukaemia; chromosome; t14; t18; q32; q21; N region; antisense; expression; inhibitor; bcl-2; IgH; immunoglobulin heavy chain; hybrid; cell death; tumour; cancer; neoplasia; diagnosis; translocation; chimera; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapies for tumours. The ribozymes may also be used to study genetic drift and mutations within cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Non-Hodgkin's B cell line bel-2/19H chimeric gene PCR primer.
                                                                                                                                                                           New enzymatic RNA molecules (ribozymes) - which cleave mRNA associated with tumours or mRNA expressed from gene encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%; Score 28; DB 14; Length 28
82.1%; Pred. No. 9.7e+03;
. wismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 28 BP; 3 A; 13 C; 7 G; 5 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1729 CCTGTGGTCCACCTGGCCCTCCGCCAAG 1756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 CCUGUGGUCCACCUGGCCCUCCGCCAAG 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CNDR ) CONSIGLIO NAZ DELLE RICERCHE,
                                                                                                                                                                                                                                                                        Claim 3; Fig 6; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT35443 standard; DNA; 30 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95IT-0MI0420.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        96WO-EP00852.
19-JAN-1993; 93US-0008910.
                                          (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-MAY-1997) (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 82.1
nes 23; Conservative
                                                                                                                                                                                                                               multiple drug resistance
                                                                                        Draper KG, Thompson JD;
                                                                                                                                   WPI; 1993-386203/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9627663-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-MAR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT35443.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT35443,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ద
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequences flanking a ACAC(CA)17 (AAQ68850) repeat element were used to design PCR primers AAQ45259 and AAQ45260. The chromosomal tocalisation was assigned to the Angelman Syndrome/Prader-Willi Syndrome region at 15q11-13. The dinuclectide repeat was confirmed to be on chromosome 15 by PCR of somatic cell hybrids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif; halrpin; hepatitis delta virus; group I intron; RNaseP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Multiple drug resistance; mdr-1; ribozyme; membrane protein; liver; resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin D; vinblastine; small intestine; kidney; adremal gland; adenocarcinoma; bowel; transformed phenotype; promyelocytic leukemia; human; chronic myelogenous leukemia; CML; follicular lymphoma; B-cell acute lymphocytic leukemia; breast cancer; colon carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                Diagnosis of genetic disorders associated with chromosomal abnormalities and uniparental disomy, e.g. Prader-Willi:Angelman syndrome - using in situ hybridisation using probes spanning the IR4-3R or GABRB3 regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cch 0.6%; Score 29.2; DB 15; 11 Similarity 91.2%; Pred. No. 6e+03; 31; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BCL-2 mRNA ribozyme cleavable nucleotide (1729).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 38 BP; 19 A; 19 C; 0 G; 0 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2306 ACATCACACACAGACAGACACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 ACACCACACACACACACACACACACACACA 34
                                                                                                                                                                                                            Ledbetter DH, Mutirangura A;
                                                                                                                                                                                                                                                                                                                                                                                                                  Example 8; Page 64; 91pp; English.
                                                                                                                                                             (BAYU ) BAYLOR COLLEGE MEDICINE
                                                                       93WO-US08501.
                                                                                                                  92US-0943639.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            93WO-0504573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92US-0882822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            92US-0882885.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92US-0936532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-0987131
93US-0006122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92US-0936422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       920S-0936531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ51954 standard; RNA; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                          WPI; 1994-118484/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
Matches 31; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                       10-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-MAY-1993;
                                                                                                                                                                                                          Airhart SD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W09323057-A.
                                                                                                                    11-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-AUG-1992
07-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JAN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-AUG-1992
                          31-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ51954;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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RESULT 16 AA051954

ð 셤 ö

Gaps

;

Length 28;

Nicolin A; Morelli S, Capaccioli S,

WPI; 1996-425424/42.

New chimeric bcl-2/IgH antisense transcript of t(14; 18) translocated cells - used to develop oligodeoxynucleotides for the diagnosis and treatment of tumours associated with the translocation.

Example 1; Page 12; 32pp; English

AAT3541-T35443 are PCR primers used to amplify the chimeric bcl-2/IgH (immunoglobulin heavy chain) gene present in the t(14: 18) chromosomally translocated non-Hodgkin's lymphoma cell line DOHH-2. The transcript produced is used for the design of antisense transcripts which can inhibit the production of bcl-2 protein in t(14: 18) translocated cells and induce cell death. The antisense transcripts can be used for the diagnosis and treatment of t(14: 18) -associated tumours and for monitoring treatment of a pathology associated with a t(14: 18) translocation 

Sequence 30 BP; 9 A; 9 C; 7 G; 5 T; 0 other;

Gaps ; 0 Score 28; DB 17; Length 30; Pred. No. 1e+04; 0; Indels 0; Mismatches Score 0.6%; s 100.0%; 28; Conservative Best Local Similarity Query Match Matches

4504 CTTCAGGGTCTTCCTGAAATGCAGTGGT 4531 δλ

30 CTTCAGGGTCTTCCTGAAATGCAGTGGT 3

용

RESULT 18

AAT35446 standard; DNA; 30 BP

AAT35446/c ID AAT354

Nen-Hodgkin's B cell line bcl-2/IgH chimeric gene PCR primer

first entry)

27-MAY-1997

Non-Hodgkin's lymphoma; cell line DOHH2; DHL-4; leukaemia; chromosome; t14; t18; q12; q11; N region; antisense; expression; inhibitor; bcl-2; lgH; immunoglobulin heavy chain; hybrid; cell death; tumour; cancer; neoplasia; diagnosis; translocation; chimera; PCR;

polymerase chain reaction; ss.

ynthetic.

WO9627663-A2 12-SEP-1996

96WO-EP00852 02-FEE 1996;

95IT-0MI0420 03-MAR-1995; CNDR ) CONSIGLIO NAZ DELLE RICERCHE.

Nicolin A; Capaccioli S, Morelli S,

WPI; 1996-425424/42.

for the New chimeric bcl-2/IgH antisense transcript of t(14; 18) translocated cells - used to develop oligodeoxynucleotides diagnosis and treatment of tumours associated with the translocation.

Example 1; Page 12; 32pp; English. 

AAT35446 and AAT35447 are PCR primers used to amplify the chimeric

bcl-2/IgH (immunoglobulin heavy chain) gene present in the t(14; 18) chromosomally translocated non-Hodgkin's lymphoma cell line DOHH-2. The primers correspond to the 3' untranslated region of the bcl-2 gene. The transcript produced is used for the design of antisense transcripts which can inhibit the production of bcl-2 protein in t(14; 18) translocated cells and induce cell death. The antisense transcripts can be used for the diagnosis and treatment of t(14; 18)-associated tumours and for monitoring treatment of a pathology associated with a t(14; 18) translocation. 

Sequence 30 BP; 9 A; 9 C; 7 G; 5 T; 0 other;

Gaps ; Length 30; 0; Indels Score 28; DB 17; Pred. No. 1e+04; 100.08; Prea. ... 0.6%; Best Local Similarity 100.0 Matches 28; Conservative

ö

4504 CTTCAGGGTCTTCCTGAAATGCAGTGGT 4531 δλ

30 CTTCAGGGTCTTCCTGAAAIGCAGTGGT g

RESULT 19 AAT35447

BP AAT35447 standard; DNA; 30

AAT35447;

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(first entry) 27-MAY-1997 Non-Hodgkin's B cell line bcl-2/IgH chimeric gene PCR primer.

Non-Hodgkin's lymphoma; cell line DOHH2; DHL-4; leukaemia; chromosome; t14; t18; q21; q21; n region; antisense; expression; infibitor; bc1-2; IgH; immunoglobulin heavy chain; hybrid; cell death; tumour; cancer; neoplasia; diagnosis; translocation; chimera; PCR; chain reaction; ss polymerase

Synthetic.

WO9627663-A2

12-SEP-1996

96WO-EP00852. 02-MAR-1996; 03-MAR-1995;

(CNDR ) CONSIGLIO NAZ DELLE RICERCHE.

Capaccioli S, Morelli S, Nicolin A;

WPI; 1996-425424/42.

New chimeric bcl-2/IgH antisense transcript of t(14; 18) translocated cells - used to develop oligodeoxynucleotides for the diagnosis and treatment of tumours associated with the translocation.

Example 1; Page 12; 32pp; English.

AAT35446 and AAT35447 are PCR primers used to amplify the chimmeric bcl-2/IgH (immunoglobulin heavy chain) gene present in the t(14; 18) chromosomally translocated non-Hodgkin's lymphoma cell line DOHH-2. The primers correspond to the 3' untranslated region of the bcl-2 gene. The transcript produced is used for the design of antisense transcripts which can inhibit the production of bcl-2 protein in t(14; 18) translocated cells and induce cell death. The antisense transcripts can be used for the diagnosis and treatment of t(14; 18)-associated tumours and for monitoring treatment of a pathology associated with a t(14; 18) translocation. 

Sequence 30 BP; 9 A; 9 C; 7 G; 5 T; 0 other;

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WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ33894/c
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                          ö
                                                                                                                                                                                            Human; Bcl-2; gene therapy; apoptosis inhibitor; mutant; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Satellite sequence; DNA fragmentation; microsatellite DNA; DNA marker; Haliotis discus; ss.
                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid sequence of Bc1-2 protein in which at least one serine residue is substituted by alanine or aspartic acid. The protein has increased apoptosis inhibitory activity compared with the wild type Bc1-2 protein. The mutated Bc1-2 protein is useful in the treatment of disorders caused by apoptosis. The present sequence was used to amplify DNA encoding the human Bc1-2 protein.
                                                                                                                                                                                                                                                                                                                                                                        New apoptosis inhibitors, useful for treating apoptosis related
                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an apoptosis inhibitor comprising the
      0.6%; Score 28; DB 17; Length 30;
100.0%; Pred. No. 1e+04;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.6%; Score 28; DB 22; I Best Local Similarity 100.0%; Pred. No. 1.1e+04; Matches 28; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 36 BP; 7 A; 10 C; 15 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1458 GATGGCGCACGCTGGGAGAACGGGGTAC 1485
                                             4504 CTTCAGGGTCTTCCTGAAATGCAGTGGT 4531
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GATGGCGCACGCTGGGAGAACGGGGTAC 36
                                                      30 CTTCAGGGTCTTCCTGAAATGCAGTGGT 3
                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 39; 43pp; Japanese.
                                                                                                                                                                         Human Bcl-2 PCR primer oligo-10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           H. discus derived sequence #11.
                                                                                                                                                                                                                                                                                                              (HISM ) HISAMITSU PHARM CO LTD.
                                                                                                                                                                                                                                                                                            99JP-0350427.
                                                                                                                                                                                                                                                                        07-DEC-2000; 2000WO-JP08667
                                                                                                                AAH45307 standard; DNA; 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ98493 standard; DNA; 40
Query Match
Best Local Similarity 100.C
"...hes 28; Conservative
                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                  Shibazaki F, Kuma H;
                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-381681/40.
                                                                                                                                                                                                                                   WO200142459-A1.
                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                            09-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JUN-2000
                                                                                                                                                      10-SEP-2001
                                                                                                                                                                                                                                                     14-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                   disorders
                                                                                                                                   AAH45307;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ98493;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 21
                                                                                              RESULT 20
                                                                                                       AAH45307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ98493
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XEX
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The invention provides a novel method for isolation of satellite sequences from genomic DNA that comprises fragmentation of the DNA by a method which is not dependent on base sequences, then selection of the satellite sequences from the obtained genomic library of high homogeneity. The method is useful for the isolation of microsatellite DNA sequences which can be used as DNA markers. The new method markedly improves the efficiency of isolation of satellite sequences in comparison to prior art methods which are reliant on base sequences. Sequences AR298483-514 represent sequences from Hallotis discus, used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolation of satellite sequences from genomic DNA for use as DNA markers comprises isolating a library with high homogeneity by DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 27.8; DB 21;
Pred. No. 1.2e+04;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 40 BP; 19 A; 19 C; 0 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                         (NORQ ) JAPAN MIN AGRIC FORESTRY & FISHERIES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Microsatellite sequence from clone TGLA309.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2309 TCACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 TCACACACACACACACACACACACACACACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 5; Page 14; 35pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAQ33894 standard; DNA; 38 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92WO-US00340.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        91US-0642342.
                                                                                                                                                                 99WO-JP03551.
                                                                                                                                                                                                                     98JP-0232153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                Takahashi H, Sekino M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-224692/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENM-) GENMARK.
Haliotis discus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fragmentation -
                                                   WO200011156-A1.
                                                                                                                                                                 01-JUL-1999;
                                                                                                                                                                                                                     18-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .5-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Simi
Matches 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-AUG-1992.
                                                                                                          02-MAR-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ33894;
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Gaps

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The invention relates to the isolation of polymorphic repeat sequences having the sequence (dd-dA)n.(dd-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis.

The repeats, when analysed, fall into 4 categories:

1) perfect repeats which are alternating tandem CA repeats with no interruptions and without adjacent repeats of another sequence.

2) imperfect repeats which are defined as 2 or more runs of uninterrupted CA repeats spearated by no more than 3 consecutive non-repeat bases;
                                                                                       containing the repeat sequences were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(dC-dA).(dG-dT) probe. Over 100 repeat blocks were isolated. The inserts from the clones were amplified by primers AAT65798-T66047. Those clones where the repeat sequence has been determined are shown in AAT65704-797. This repeat sequence is from the marker clone Mdf42 which contains the repeat
having the sequence (dC-dA)n.(dG-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis. Clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Polymorphism; repeat sequence; genetic marker; primer; amplification; PCR; polymerase chain reaction; paternity; maternity; human; pedigree; linkage analysis; genetic disease; animal; plant; breeding; locus; hybridisation; chromosome; ds.
                                                                                                                                                                                                                                                                             0.5%; Score 27.2; DB 18; Length 40; 80.0%; Pred. No. 1.7e+04; ive 0; Mismatches 8; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n - using novel nucleic acid mols. as primers
                                                                                                                                                                                                                                                                                                                                                          2310 CACACACACAGACACACACACACACACAACAATTAACA 2349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (dC-dA)n.(dG-dT)n polymorphic repeat sequence #3.
                                                                                                                                                                                                                                                                                                                                                                            Sequence 40 BP; 20 A; 19 C; 0 G; 1 T; 0 other;
                                                                                                                                                                                                          the formula: (CA)16T(AC)3.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 8; Column 57-58; 186pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT66054 standard; DNA; 40 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91US-0754351.
89US-0341562.
94US-0222177.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MARS-) MARSHFIELD CLINIC.
                                                                                                                                                                                                                                                                                                Local Similarity 80.0 les 32; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1997-042299/04
                                                                                                                                                                                                          sequence having
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-APR-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US5582979-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-APR-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT66054;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Weber JL;
                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT66054
                                                                                                                                                                                                                                                                                                                                                          ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22222222222228
                                                                                                     The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine MboI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonoucleotide probe.
Cone out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (TG)n >9 microsatellites in the bovine genome is estimated at >100, 000. The sequence information for ac. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). The sequences upstream and downstream of the microsatellite sequence were used to generate the required PCR primmers for in vitro amplification of the corresp. microsatellite (using the program of the microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait closi, or genes involved the determinism of economically important set is also AAQ33501-34437.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphism; repeat sequence; genetic marker; primer; amplification; PCR; polymerase chain reaction; paternity; maternity; human; pedigree; linkage analysis; genetic disease; animal; plant; breeding; locus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to the isolation of polymorphic repeat sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                       Polymorphic bovine DNA markers - used in genetic identification,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n - using novel nucleic acid mols. as primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Repeat sequence from polymorphic marker clone Mfd42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.5%; Score 27.6; DB 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 1.3e+04;
0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 38 BP; 1 A; 0 C; 19 G; 18 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2306 ACATCACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       37 ACAACACACACACACACACACACACTCACACA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Column 9-10; 186pp; English.
                                      gene mapping, and selective breeding
                                                                            Table 7; Page 287; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hybridisation; chromosome; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT65743 standard; DNA; 40 BP
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94US-0222177.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91US-0754351.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JUN-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MARS-) MARSHFIELD CLINIC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best_Local Similarity 88.2
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1997-042299/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-APR-1989;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-APR-1989;
04-APR-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 23
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3) compound perfect repeats which are uninterrupted runs of CA separated by no more than 3 consecutive non-repeat bases from a run of at least 5 uninterrupted dinuclectide or longer repeats of a sequence other than (dC-dA)n. (dG-dT)n, or from at least 10 uninterrupted mononuclectides; and 4) imperfect compound repeats which are defined as for the perfect compound repeats except that the runs of CA are interrupted. The sequence presented here is an example of a perfect repeat sequence of structure: (CA)20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a substantially purified or isolated nucleic acid (I) from ryegrass or fescue species including a simple sequence repeat (SSR), having 2 or more tandemly repeated nucleotide core elements 2-6 nucleotides in length. Also included are a nucleic acid primer suitable for amplifying an SSR, identifying (M1) an SSR by preparing a library of ryegrass or fescue genomic DNA enriched for SSRs and identifying clones in the library containing SSRs, a library of ryegrass or fescue genomic DNA enriched for SSRs prepared by the M1, selecting for a gene in grass or cereal breeding by identifying an SSR that is closely associated with the gene such that the SSR and the gene are preferentially co-inherited, and selecting for the SSR in the breeding, a method for DNA profiling grass or cereal species varieties by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Simple sequence repeat; plant; ds; SSR; ryegrass; fescue; tandem repeat; cereal profiling; grass profiling; seed batch purity testing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New simple sequence repeats having 2 or more tandemly repeated nucleotide core elements isolated from ryegrass and fescue, useful for selecting of genes in grass or cereal breeding or profiling grass or
                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                           .;
0
                                                                                                                                                                                                                 Score 27.2; DB 18; Length 40; Pred. No. 1.7e+04; 0; Mismatches 8; Indels (
                                                                                                                                                                                                                                                                                           2310 CACACACACAGACAGACACACACACACAATTAACA 2349
                                                                                                                                                                                                                                                                                                               1 CACACACACACACACACACACACACACACACA 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SAUS-) STATE SOUTH AUSTRALIA SOUTH AUSTRALIAN R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         UNIV SOUTHERN CROSS.
STATE VICTORIA DEPT NATURAL RES & ENVIRO.
                                                                                                                                                                           Sequence 40 BP; 20 A; 20 C; 0 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ITMA-) INT MAIZE & WHEAT IMPROVEMENT CENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Simple sequence repeat, SSR, #44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 13; Page 53; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                AAS13773 standard; DNA; 40 BP
                                                                                                                                                                                                                   0.5%;
                                                                                                                                                                                                                                   80.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-JAN-2001; 2001NZ-0509193.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99AU-0004906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-MAY-2000; 2000AU-0007310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                   Local Similarity 80.0 nes 32; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cereal species varieties
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jones ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UNIV ADELAIDE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-512563/56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NZ509193-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS13773;
                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYSC-)
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                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                          RESULT 25
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The invention relates to a method of identifying or detecting a nucleotide repeat region in a nucleic acid molecule characterised by a particular length, comprising employing ligase-assisted spacer additton (LASA) assay. The method is useful in the identifying or detecting a nucleotide repeat region in a nucleic acid molecule characterised by a particular length. In particular, the method is useful for identification of a nucleotide length polymorphism in animals or humans, which is associated with a neurodegenerative disease including fragile X sociated with a neurodegenerative disease including fragile X pracrome, Huntington's disease, or muscular dystrophy. Furthermore, the method may be used for identifying and/or typing microorganisms including yeasts and lower uni- and multi-callular organisms, as well as prokaryotic microorganisms; and for genotyping subjects including humans. The method is also useful for detecting certain cancers and other malkers for use in identification of human and non-human individuals, plants and
                                                                                                                                                                                                                                           ö
assessing variation between SSR varieties and testing the purity of great or cereal seed batches by assessing variation within seed batch of an SSR. The SSRs may be used in the selection of genes in grass or cereal breeding, for profiling grass or cereal species varieties, for testing the purity of grass or cereal seed batches, and for DNA profiling to establish the distinct identity, uniformity and/or stability of a cultivar. The present sequence is a ryegrass or fescue SSR.
                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Microsatellite; ligase-assisted spacer addition assay; LASA; cancer; nucleotide length polymorphism detection; neurodegenerative disease; fragile X syndrome; Huntington's disease; muscular dystrophy; forensic; gene mapping; population study; human; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphism associated with a neurodegenerative disease, comprises using a ligase-assisted spacer addition assay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting a nucleotide repeat region in a nucleic acid having a particular length, useful for identifying nucleotide length
                                                                                                                                                                                                                                           ;
0
                                                                                                                                                                                                    Length 40;
                                                                                                                                                                                                                                           8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human microsatellite D1S191 detection PCR primer #8.
                                                                                                                                                                                                                                                                               2310 CACACACACAGACAGACACACACACACAACAATTAACA 2349
                                                                                                                                                                                                    Score 27.2; DB 23;
Pred. No. 1.7e+04;
                                                                                                                                                                                                                                                                                                      Barnard R,
                                                                                                                                                            Sequence 40 BP; 20 A; 20 C; 0 G; 0 U; 0 other;
                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wolter L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10; Page 55; 89pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                         ABK24300 standard; DNA; 40 BP
                                                                                                                                                                                                    0.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      39-MAY-2001; 2001WO-AU00526.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-MAY-2000; 2000US-202771P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DIAT-) DIATECH PTY LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brockhurst V, Timms P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-121948/16.
                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                      Ma.
Local 5..
32;
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                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                           Matches
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microorganisms, to ascertain parentage of human or non-human individual, and to monitor responses to therapies including the possibility of uncleic acid damage. The nucleotide polymorphisms may be used in forensic science to identify a particular victim or an alleged perpetrator of a crime, in gene mapping and population studies. LASA may also be used in the manufacture of a kit for detecting and/or identifying nucleotide repeat regions such as a nucleotide length polymorphism in a eukaryotic genome. The LASA method avoids the time and cost required by prior art methods using gel electrophoresis and Southern transfer analysis.

The particular, current diagnosis of Huntington's disease relies heavily upon the use of gel electrophoresis, a process that has proved difficult to automate or miniaturise. The LASA method allows total confidence of this limiting step, making it a strong candidate for future use in clinical and laboratory procedures. ABK24276-ABK24313 represent primers used to detect polymorphisms or microsatellites as described in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Detecting target sequences including chromosomal abnormality - is by amplification then using separable probe and detectable probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        detection of chromosomal abnormalities; translocations; neoplasm; cancer; follicular lymphoma; T lymphocytes; B lymphocytes;
                                                                                                                                                                                                                                                                                                                                       ;
0
                                                                                                                                                                                                                                                                                                        Score 27.2; DB 24; Length 40;
                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Turchetti E;
                                                                                                                                                                                                                                                                                                                                                                      2310 CACACACACAGACAGACACACACACACAACAATTAACA 2349
                                                                                                                                                                                                                                                                                                                       Pred. No. 1.7e+04;
0; Mismatches 8;
                                                                                                                                                                                                                                                                                                                                                                                         Sequence 40 BP; 20 A; 20 C; 0 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Martinazzo G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 22; 28pp; English
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/note= "amidated"
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"amidated"
                                                                                                                                                                                                                                                                                                          0.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            91IT-0000286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAQ30102 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-MAR-1993 (first entry)
                                                                                                                                                                                                                                                                                                                        Best_Local Similarity 80.0
Matches 32; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Marcolini S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MBR reporter sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1992-398894/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           high specificity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-APR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W09219775-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ30102;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bichi R,
                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ30102
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This reporter is used to determine which breakpoint is present in the MBR and mcr amplified sequences detected using AAQ30096-101.

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The sequences given in AAQ51825-2266 represent areas of mRNAs which are associated with development or maintenance of chronic myelogenous leukemia (CML), promyelocytic leukemia, Burkitt's lymphoma, or acute lymphocytic leukemia, follicular lymphoma, B-cell acute
The terminal amino gps. are used to conjugate the reporter to alkaline phosphatase, making them "signal generating" probes. They are used in a method for the detection of chromosomal abnormalities eg. translocations correlated to a neoplastic state such as those related to T and B lymphocytes eg. follicular lymphoma. The method provides a quantitative measurement and provides a veryhigh degree of specificity minimising the risk of false positives.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neuroblastoma; lung cancer; genetic drift; mutation; hammerhead motif; hairpin; hepatitis delta virus; group I intron; RNaseP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Multiple drug resistance; mdr-1; ribozyme; membrane protein; liver; resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin D; vinblastine; small intestine; kidney; adrenal gland; adenocarcinoma; bowel; transformed phenotype; promyelocytic leukemia; human; chronic myelogenous leukemia; CML; follicular lymphoma; B-cell acute lymphocytic leukemia; breast cancer; colon carcinoma;
                                                                                                                                                                                                 Gaps
                                                                                                                                                                                               ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New enzymatic RNA molecules (ribozymes) - which cleave mRNA associated with tumours or mRNA expressed from gene encoding
                                                                                                                                                               Length 27;
                                                                                                                                                                                               0; Indels
                                                                                                                                                             Score 27; DB 13; I
Pred. No. 1.6e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                           BCL-2 mRNA ribozyme cleavable nucleotide (2081).
                                                                                                                                 Sequence 27 BP; 8 A; 12 C; 3 G; 4 T; 0 other;
                                                                                                                                                     0.5%; Scor.
100.0%; Pred. No. ...
0; Mismatches
                                                                                                                                                                                                                                4440 IIICAACACAGACCCACCCAGAGCCCI 4466
                                                                                                                                                                                                                                                  1 TTTCAACACAGACCCACCAGAGCCCT 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Fig 6; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                AAQ51963 standard; RNA; 27 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      920S-0936421.
920S-0936422.
920S-0936531.
920S-0936532.
920S-0987131.
930S-0006122.
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92US-0936110
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                   27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      multiple drug resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Draper KG, Thompson JD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1993-386203/48.
                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                              26-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9323057-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-JAN-1993
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                                                                                                                                                                                                                                                                                                                                                                                AAQ51963;
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                                                                                                                                                                                                                                                                                                                 RESULT 28
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Gaps

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0; Indels

Mismatches

1970 ACAACATCGCCCTGTGGATGACTGAGT 1996

0.5%; Score 27; 100.0%; Pred. No. tive 0; Mismatch

Query Match 0.5 Best Local Similarity 100. Matches 27; Conservative

Length 27;

DB 24; L

Sequence 27 BP; 6 A; 7 C; 7 G; 7 T; 0 other;

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lymphocytic leukemia, breast cancer, colon carcinoma, neuroblastoma and lung cancer. The full length mRNAs containing these target sequences, encode aberant cellular proteins which are able to control cellular proliferation and are directly linked to a leukemic phenotype. These target sequences are identified by the ribozyme of the invention. The ribozymes is formed in a hammerhead motiff, but may also be formed in the motif of a hairpin, hepatitis dolta virus, group intron or RNaseP-like RNA. These ribozymes may be used to inhibit the development or expression of a transformed phenotype in man and other animals by modulating expression of the corresponding gene. Cleavage of target mRNAs expressed in pre-neoplastic and transformed cells elicits inhibition of the transformed state. Multiple drug cells elicits inhibition of the transformed state. Multiple drug drug resistance used by transformed cells and thus enhances drug the resistance used by transformed cells and thus enhances drug the resistance used by transformed cells and thus enhances drug the resistance used by transformed cells and thus enhances drug the resistance used by transformed cells and thus enhances drug the resistance used by transformed cells and thus enhances drug the resistance used by transformed cells and thus enhances drug the modulation of the ribozymes remove the mechanism of the removed the recent of the removed the mechanism of the removed removed removed the removed rem
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            therapies for tumours. The ribozymes may genetic drift and mutations within cells.
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Sequence 27 BP; 1 A; 8 C; 5 G; 13 U; 0 other;

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Gaps
                                            ;
  Length 27;
  ch 0.5%; Score 27; DB 14; Length 27 il Similarity 51.9%; Pred. No. 1.6e+04; 14; Conservative 13; Mismatches 0; Indels
Query Match
Best Local Similarity
Matches 14; Conserv
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2081 CTCTGTTTGATTTCTCCTGGCTGTCTC 2107 

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AAL46663 standard; DNA; 27 Human bcl-2 mRNA probe. 05-AUG-2002 AAL46663; RESULT 29 AAL46663/c 

(first entry)

Human; bcl-2; cancer detection; disseminated cancer cell; cytostatic; probe; ss.

Homo sapiens.

WO200237113-A2.

10-MAY-2002.

J5-NOV-2001; 2001WO-EP12786.

03-NOV-2000; 2000DE-1054635.

(GIES/) GIESING M.

Boeckmann B, Suchy B; Siesing M, Grill H, 

WPI; 2002-426739/45.

Clinically validating target from disseminated cancer cells by determining whether status of target determined in cancer cells of individuals correlates with cancer-related information about clinical status of individuals -

Example 3; Page 55; 57pp; English.

a target from disseminated cancer cells, characterised in that for a population of individuals it is determined whether a status of the target determined in disseminated cancer cells of the individuals correlates with at least one cancer-related information about the clinical status of the individuals. The method is useful for clinically validating target from disseminated cancer cells. The present sequence is a probe used to demonstrate the method of the invention. The present invention relates to a method for the clinical validation of

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine MoI DNA fragments of between 250 and 500 bp with an (AC)15 and a (FC)15 oligonucleotide probe. One out of 50 chones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (FG)n >9 microsatellites in the bovine genome is estimated at >100, 000. The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The sequences upstream and downstream of the microsatellite sequence were used to generate the required PCR primers for in vitro amplification of the corresp. microsatellite sequence (Using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait traits esp. in cattle, to allow selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polymorphic bovine DNA markers - used in genetic identification,
                                                                                                                                                                                                                                PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.5%; Score 27; DB 13; Length 39 Best Local Similarity 85.7%; Pred. No. 1.8e+04; Matches 30; Conservative 0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 39 BP; 2 A; 0 C; 17 G; 20 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2310 CACACACACAGACAGACACACACACACAACAAT 2344
                                                                                                                                                                                              Microsatellite sequence from clone TGLA153.
                                                                                                                                                                                                                                                genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene mapping, and selective breeding
27 ACAACATCGCCCTGTGGATGACTGAGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Table 7; Page 225; 517pp; English.
                                                                                              AAQ33737 standard; DNA; 39 BP
                                                                                                                                                                                                                                                                                                                                                                                 92WO-US00340.
                                                                                                                                                                                                                                                                                                                                                                                                                  91US-0642342.
                                                                                                                                                              02-FEB-1993 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENM-) GENMARK.
                                                                                                                                                                                                                                                                                                                                                                                 15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                  15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                  06-AUG-1992.
                                                                                                                                                                                                                                                                                                                 WO9213102-A.
                                                                                                                                                                                                                                                                                 Bos taurus.
                                                                                                                               AAQ33737;
                                                              RESULT 30
AAQ33737/c
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PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage;
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                                                                            25-MAY-2001.
                                                     NZ509193-A.
                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ34032;
                                                                                                                                                                                   (VICT-)
                                                                                                                                                                                            UYAD-)
                                                                                                                                                                        UYSC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 33
AAQ34032/c
δλ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         XX EX EX EX SX
                                                                                                                                                                                                                                                                                                                                                                                                 The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine Mooi DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe.
250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe.
250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe.
251 one out of 50 clones cross-hybridised. Assuming independent of distribution of microsatellites and Mbol sites, the frequency of (TC) >9 microsatellites and Mbol sites, the frequency of (TC) >9 microsatellites in the bovine genome is estimated at >100, construction of the specification and indexed herein (see below).
252 The sequences upstream and downstream of the microsatellite sequence were used to generate the required PCR primmers for in vitro amplification of the corresp. microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, corresp. in cattle, the determinism of economic trait clori, or genes involved the determinism of economically important craits esp. in cattle, to allow selective breeding.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                Polymorphic bovine DNA markers - used in genetic identification, gene mapping, and selective breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ó;
                                                                                                                            selection; primers; OPTIPRIM; breeding; cattle; parentage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.5%; Score 26.8; DB 13;
93.3%; Pred. No. 1.8e+04;
ative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30 BP; 0 A; 0 C; 15 G; 15 T; 0 other;
                                                                                                      Microsatellite sequence from clone TGLA357.
                                                                                                                                        genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30 CACACACACACACACACACACACACACA 1
                                                                                                                                                                                                                                                                                                                                                                                   rable 7; Page 316; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Simple sequence repeat, SSR, #59.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAS13788 standard; DNA; 30 BP
                                    AA033965 standard, DNA; 30 BP.
                                                                                                                                                                                                                                 92WO-US00340.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 93.3
nes 28; Conservative
                                                                                                                                                                                                                                                                                                                           WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                               (GENM-) GENMARK.
                                                                                                                                                                                                                                                        15-JAN-1991;
                                                                                02-FEB-1993
                                                                                                                                                                                                            06-AUG-1992
                                                                                                                                                                                     WO9213102-A
                                                                                                                                                                                                                                                                                                       Georges M,
                                                                                                                                                                Bos taurus.
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                                                         AAQ33965;
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                         AAQ33965/c
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               RESULT 31
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The invention relates to a substantially purified or isolated nucleic acid (1) from ryegrass or fescue species including a simple sequence repeat (SSR), having 2 or more tandemly repeated nucleotide core elements core included are a nucleic acid primer prince and primers of ryegrass or fescue genomic DRA enriched for SRRs and included are a nucleic acid primer of ryegrass or fescue genomic DRA enriched for SRRs and included in the library containing SSRs, a library of ryegrass or fescue genomic DRA enriched for SRRs prepared by the M1, selecting for a gene in grass or cereal breeding by identifying an SSR that is closely cassociated with the gene such that the SSR in the selecting for the SSR in the second backed for DNA profiling grass or cereal species varieties by assessing variation within seed batch of an SSR. The SSRs may be used in the selection of genes in grass or cereal core breeding, for profiling grass or cereal species varieties, for testing the purity of grass or cereal seed batches, and for DNA profiling to cetablish the distinct identity, uniformity and/or stability of a cultivar. The present sequence is a ryegrass or fescue SSR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
Simple sequence repeat; plant; ds; SSR; ryegrass; fescue; tandem repeat; cereal profiling; grass profiling; seed batch purity testing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New simple sequence repeats having 2 or more tandemly repeated nucleotide core elements isolated from ryegrass and fescue, useful for selecting of genes in grass or cereal breeding or profiling grass or cereal species varieties
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      UNIV SOUTHERN CROSS.
STATE VICTORIA DEPT NATURAL RES & ENVIRO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE SOUTH AUSTRALIA SOUTH AUSTRALIAN R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 30 BP; 15 A; 15 C; 0 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Microsatellite sequence from clone TGLA429.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2310 CACACACAGACAGACACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 13; Page 53; 72pp; English.
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                                                                                                                                                                                                                                                                                                                                            03-JAN-2001; 2001NZ-0509193.
                                                                                                                                                                                                                                                                                                                                                                                                                       24-DEC-1999; 99AU-0004906.
04-MAY-2000; 2000AU-0007310.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Forster JW, Jones ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UNIV ADELAIDE
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The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine Mool DNA fragments of between 250 and 500 bp with an (AC)15 and a (FC)15 oligonucleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and Mbol sites, the frequency of (FG)n >9 microsatellites in the bovine genome is estimated at >100, 000. The sequence information for ca. 230 such bovine microsatellites in the specification and indexed herein (see below).
                                                                                                                                                                                                                                                                                                                                                                                                                            The sequences upstream and downstream of the microsatellite sequence were used to generate the required PCR primers for in vitro amplification of the corresp. microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding.
                                                                                                                                                        Polymorphic bovine DNA markers - used in genetic identification,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 26.8; DB 13;
Pred. No. 1.9e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 31 BP; 0 A; 0 C; 15 G; 16 T; 0 other;
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                                                                                                                                                                               gene mapping, and selective breeding
                                                                                                                                                                                                                  Table 7; Page 349; 517pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                See also AAQ33501-34437.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genetic mapping; traits;
                                                                         Georges M, Massey JM;
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                                                                                                            WPI; 1992-284684/34.
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                               (GENM-) GENMARK.
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Cone out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (TC)n >9 microsatellites in the bovine genome is estimated at >100, co. The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). The sequence upstream and downstream of the microsatellite sequence were used to generate the required PCR primars for in vitro amplification of the corresp. microsatellite (using the program OPTIRRIM). The microsatellites may be used to identify individuals, correspecting, and in the genetic mapping of economic trait closi, or genes invalved the determinism of economic trait traits esp in cattle, to allow selective breeding.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 31 BP; 0 A; 0 C; 15 G; 16 T; 0 other;
genetic mapping; traits; amplification; ss.
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                                                                                                                                                                                                                                                                                                  Georges M, Massey JM;
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                                                                                                                                                                                                                                                      (GENM-) GENMARK
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                                          Bos taurus.
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NAMES OF STREET OF STREET

RESULT 34

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Length 31; Indels

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The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine MboI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe.

CC 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe.

CC one out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (TG)n >9 microsatellites in the bovine genome is estimated at >100, 200 000. The sequence information for a. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below).

CT we sequence upstream and downstream of the microsatellite sequence used to generate the required PCR primmers for in vitro amplification of the corresp. microsatellite (using the program optimization). The microsatellites may be used to identify individuals, and in the genetic mapping of economic trait clori, or genes involved the determinism of economically important traits sep. in cattle, to allow selective breeding.
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                                                                                                                                                                                                                                                                                                                                                                                 0.5%; Score 26.8; DB 13; Length 31; 93.3%; Pred. No. 1.9e+04; Live 0; Mismatches 2; Indels C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genetic mapping; traits; amplification; ss
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                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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AAQ33822/c
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screening a library of bovine MDOI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MDOI sites, the frequency of (T6)n >9 microsatellites in the bovine genome is estimated at >100, The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            is summarised in the specification and indexed herein (see below). The sequences upstream and downstream of the microsatellite sequence
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amplification of the corresp. microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding.

See also AAQ33501-34437.
                                                                                                                                                                                          Gaps
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                                                                                                                                                      0.5%; Score 26.8; .DB 13; Length 31; 93.3%; Pred. No. 1.9e+04; tive 0; Mismatches 2; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    traits esp. in cattle, to allow selective breeding.
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Pred. No. 1.9e+04;
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                                                                                                                       Sequence 31 BP; 0 A; 0 C; 15 G; 16 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  genetic mapping; traits; amplification; ss.
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                                                                                                                                                                               Local Similarity 93.33
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Best Local Similarity
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AAT65759
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n.(dG-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis. Clones chromosome-specific phage untrained by hybridisation of chromosome-specific phage ilbraries with a synthetic poly(dC-dA). (dG-dT) probe. Over 100 repeat blocks were isolated by hybridisation of clones were amplified by primers Arafo799 # T6047. Those clones where the repeat sequence has been determined are shown in AAT65704-797. This repeat sequence is from the marker clone Mdf79 which contains the repeat sequence having the formula: (AC)15.5.
                                                                                                                                                                                Polymorphism; repeat sequence; genetic marker; primer; amplification; PCR; polymerase chain reaction; paternity; maternity; human; pedigree; linkage analysis; genetic disease; animal; plant; breeding; locus;
       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                          Repeat sequence from polymorphic marker clone Mfd79.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 31 BP; 16 A; 15 C; 0 G; 0 U; 0 other;
   0; Mismatches
                   2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Column 11-12; 186pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2310 CACACACAGACAGACACACACACACACA 2339
                               31 CACACACACACACACACACACACACACACA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 CACACACACACACACACACACACACACA 31
                                                                                                                                                                                                                 hybridisation; chromosome; ds.
                                                                                                 BP.
                                                                                                                                                                                                                                                                                                                        91US-0754351.
89US-0341562.
94US-0222177.
                                                                                                                                                                                                                                                                                                      89US-0341562,
                                                                                             AAT65765 standard; DNA; 31
                                                                                                                                     17-JUN-1997 (first entry)
 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                (MARS-) MARSHFIELD CLINIC.
                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-042299/04.
                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                         US5582979-A.
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                                                                                                                                                                                                                                                                                                                       05-SEP-1991;
                                                                                                                                                                                                                                                                                                                                             04-APR-1994;
                                                                                                                 AAT65765;
                                                                                                                                                                                                                                                                                                                                                                                      Weber JL;
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Matches
                                                                        RESULT 38
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                                                                                   AAT65765
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The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n.(dG-dT)n which can be used as genetic markers. Primars based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis. Clones containing the repeat sequences were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(dC-dA).(dG-dT) probe. Over 100 repeat blocks were isolated. The inserts from the clones were amplified by primers AAR5798-766047. Those clones where the repeat sequence has been determined are shown in AAR55704-797. This sequence having the formula: (CA)15.5.
                                                                                                       Polymorphism; repeat sequence; genetic marker; primer; amplification; PCR; polymerase chain reaction; paternity; maternity; human; pedigree; linkage analysis; genetic disease; animal; plant; breeding; locus; hybridisation; chromosome; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n - using novel nucleic acid mols. as primers
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                                                         Repeat sequence from polymorphic marker clone Mfd57.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.5%; Score 26.8; DB 18;
13.3%; Pred. No. 1.9e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 31 BP; 15 A; 16 C; 0 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Column 11-12; 186pp; English.
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89US-0341562.
94US-022177.
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les 28; Conservative
17-JUN-1997 (first entry)
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                                                                                                                                                                                                                                                 Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-APR-1994;
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AAT65753 standard; DNA; 31 BP.

AAT65753 ID AAT6

Weber JL;

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The sequence is that of a bovine microsatellite sequence obtd. by screening a library of bovine MboI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 algonucleotide probe. One out of 50 clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (T6)n >9 microsatellites in the bovine genome is estimated at >10, co. 000. The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). The sequences upstream and downstream of the microsatellite sequence were used to generate the required PCN primers for in vitro amplification of the corresp. microsatellite (using the program OPTIREM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait coi, or genes involved the determinism of economically important criaits esp. in cattle, to allow selective breeding.
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                                                                                                                      Polymorphic bovine DNA markers - used in genetic identification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Repeat sequence from polymorphic marker clone Mfd119.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       / Match 0.5%; Score 26.8; DB 13; Local Similarity 93.3%; Pred. No. 1.9e+04; les 28; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 32 BP; 0 A; 0 C; 16 G; 16 T; 0 other;
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                                                                                                                                                gene mapping, and selective breeding
                                                                                                                                                                                          rable 7; Page 196; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT65790 standard; DNA; 32 BP
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89US-0341562.
94US-0222177.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        See also AAQ33501-34437.
                                     Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-042299/04.
                                                                                   WPI; 1992-284684/34.
(GENM-) GENMARK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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21-APR-1989;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              qq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to the isolation of polymorphic repeat sequences having the sequence (dc-dh)n.(dg-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis. Clones containing the repeat sequences were isolated by hybridisation of probe. Over 100 repeat blocks were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(dc-dh).(dg-dT) probe. Over 100 repeat blocks were isolated. The inserts from the clones were amplified by primers AAr65798 r66047. Those clones where the repeat sequence has been determined are shown in AAr65704-797. This repeat sequence is from the marker clone Mdf64 which contains the repeat sequence having the formula: (AC)15.5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
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93.3%; Pred. No. 1.9e+04;
ative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 31 BP; 16 A; 15 C; 0 G; 0 U; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Column 11-12; 186pp; English.
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    hybridisation; chromosome; ds
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94US-0222177.
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                                                                                                                                                                                                                                                                                                               (MARS-) MARSHFIELD CLINIC
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                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-042299/04.
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                                                Homo sapiens.
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                                                                                        US5582979-A.
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Query Match

Matches

ò 셤 AAQ33666;

RESULT 41 AAQ33666/c

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Gaps

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The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n.(dG-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis. Clones containing the repeat sequences were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(dC-dA). (dG-dT) probe. Over 100 repeat blocks were isolated. The inserts from the clones were amplified by primers AAPG5798-T66047. Those clones where the repeat sequence has been determined are shown in AAPG5704-797. This repeat sequence is from the marker clone Mdfill9 which contains the repeat sequence having the formula: (AC)16.
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(dC-dA)n(dG-dT)n - using novel nucleic acid mols, as primers
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Local Similarity 93.3%; Pred. No. 1.9e+04;
Local Similarity 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 32 BP; 16 A; 16 C; 0 G; 0 U; 0 other;
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                                                                        Claim 1; Column 13-14; 186pp; English.
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94US-0222177.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MARS-) MARSHFIELD CLINIC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1997-042299/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-APR-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US5582979-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-DEC-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Weber JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 43
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containing the repeat sequences were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(dC-dA).(dG-dT) probe. Over 100 repeat blocks were isolated. The inserts from the clones were amplified by primers AAT05798-766047. Those clones where the repeat sequence has been determined are shown in AA065704-797. This repeat sequence is from the marker clone Mdf55 which contains the repeat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; alpha-7 nicotinic receptor; neuronal; hybridisation; probe;
alpha-7 neuronal nicotinic acetylcholine receptor; schizophrenia;
small cell lung carcinoma; breast cancer; nicotine-dependent illness;
epilepsy; juvenile myoclonic epilepsy; Prader-Willi syndrome;
                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human alpha-7 neuronal nicotinic acetylcholine receptor and related
                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                         Length 32;
                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human alpha-7 nicotinic receptor probe SEQ ID NO:10.
                                                                                                                                                                                                     0.5%; Score 26.8; DB 18;
33.3%; Pred. No. 1.9e+04;
ve 0; Mismatches 2;
                                                                                                                                                              Sequence 32 BP; 16 A; 16 C; 0 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                  2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                       2 CACACACACACACACACACACACACACACA 31
                                                                                                                      sequence having the formula: (AC)16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Angelman's syndrome; PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 9; Page 63; 104pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                       AAX56155 standard; DNA; 32 BP
                                                                                                                                                                                                                        93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98WO-US21762.
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                                                                                                                                                                                                                                             28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Freedman R, Leonard S;
                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-288306/24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (FREE/) FREEDMAN R. (LEON/) LEONARD S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-OCT-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX56155;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                  RESULT 44
                                                                                                                                                                                                                                                                                                                                                                                                       AAX56155
                                                                                                                                                                                                                            Best
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Query Match
             ABK24296;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention provides a novel method for isolation of satellite sequences from genomic DNA that comprises fragmentation of the DNA by a method which is not dependent on base sequences, then selection of the satellite sequences from the obtained genomic library of high homogeneity. The method is useful for the isolation of microsatellite DNA sequences which can be used as DNA markers. The new method markedly improves the efficiency of isolation of satellite sequences in comparison to prior art methods which are reliant on base sequences. Sequences AA298483-514 represent sequences from Haliotis discus, used in the method of the invention.
                                                                                                                                                                                                                                                                                                                 Satellite sequence; DNA fragmentation; microsatellite DNA; DNA marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolation of satellite sequences from genomic DNA for use as DNA markers comprises isolating a library with high homogeneity by DNA fragmentation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.5%; Score 26.8; DB 21; Length 32; 93.3%; Pred. No. 1.9e+04; tive 0; Mismatches 2; Indels (
                                                      Length 32;
                                                                                  Indels
  epilepsy) and Prader-Willi and Angelman's syndromes.
                                                     0.5%; Score 26.8; DB 20;
Local Similarity 93.3%; Pred. No. 1.9e+04;
les 28; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 32 BP; 16 A; 16 C; 0 G; 0 U; 0 other;
                            Sequence 32 BP; 16 A; 16 C; 0 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NORQ ) JAPAN MIN AGRIC FORESTRY & FISHERIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. 1.96
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                            2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 CACACACACACACACACACACACACACA 30
                                                                                                                             1 CACACACACACACACACACACACACACACACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 5; Page 14; 35pp; Japanese.
                                                                                                                                                                                                        AAZ98487 standard; DNA; 32 BP
                                                                                                                                                                                                                                                                                            H. discus derived sequence #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-JP03551.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98JP-0232153.
                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 93.3 hes 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pakahashi H, Sekino M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-224692/19.
                                                                                                                                                                                                                                                                                                                                     Haliotis discus; ss.
                                                                                                                                                                                                                                                                                                                                                               Haliotis discus.
                                                                                                                                                                                                                                                                                                                                                                                         WO200011156-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-AUG-1998;
                                                                                                                                                                                                                                                                 19-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                    02-MAR-2000
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                                                                                                                                                                                                                                       AAZ98487;
                                                         Query Match
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                                                                                     Matches
                                                                                                                                                                                 RESULT 45
                                                                                                                                                                                                AAZ98487
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The invention relates to a method of identifying or detecting a nucleotide repeat region in a nucleic acid molecule characterised by a particular length, comprising employing ligase-assisted spacer addition (LASA) assay. The method is useful in the identifying or detecting a nucleotide repeat region in a nucleic acid molecule characterised by a nucleotide repeat region in a nucleic acid molecule characterised by a particular length. In particular, the method is useful for identification of a nucleotide length polymorphism in animals or humans, which is associated with a neurodegenerative disease including fragile X associated with a neurodegenerative disease including fragile X expurome, Huntington's disease, or muscular dystrophy. Furthermore, the method is also used for identifying and/or typing microorganisms including yeasts and lower uni- and multi-cellular organisms, as well as prokaryotic microorganisms; and for genotyping subjects including humans. The method is also useful for detecting certain cancers and other malignancies. Moreover, the method can be used to provide markers for use in identification of human and non-human individuals, plants and confurnant or monitor responses to therapies including the possibility of and to monitor responses to therapies including the possibility of a crime, in gene mapping and population studies. LASA may also be used in the manufacture of a kit for detecting and/or identifying nucleotide crime if he LASA method avoids the time and cost required by prior art methods using mel alength polymorphism in a eukaryotic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              art methods using generation of the process and Southern transfer analysis. In particular, current diagnosis of Huntington's disease relies heavily upon the use of gel electrophoresis, a process that has proved difficult to automate or miniaturise. The LASA method allows total avoidance of this limiting step, making it a strong candidate for future use in clinical and laboratory procedures. ABK24276-ABK24313 represent primers used to detect polymorphisms or microsatellites as described in the method of the invention.
                                                                                                                                                                         Microsatellite; ligase-assisted spacer addition assay; LASA; cancer; nuclectide length polymorphism detection; neurodegenerative disease; fragile X syndrome; Huntington's disease; muscular dystrophy; forensic; gene mapping; population study; human; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detecting a nucleotide repeat region in a nucleic acid having a particular length, useful for identifying nucleotide length polymorphism associated with a neurodegenerative disease, comprises using a ligase-assisted spacer addition assay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Giffard PM;
                                                                                       Human microsatellite D1S191 detection PCR primer #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Brockhurst V, Timms P, Wolter L, Barnard R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 10; Page 55; 89pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-MAY-2001; 2001WO-AU00526.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-MAY-2000; 2000US-202771P.
10-MAY-2000; 2000US-202559P.
39-APR-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
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Gaps

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Indels

0; Mismatches

Local Similarity 93.3%; les 28; Conservative

Matches

ABK24296 standard; DNA; 32 BP

RESULT 46 ABK24296

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0.5%; Score 26.8; DB 24; Length 32; 13.3%; Pred. No. 1.9e+04;

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AAT65705;
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NA PROPERTY OF THE PROPERTY OF
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250 and 500 bp with an (AC)15 and a (TC)15 oligonuclectide probe.

260 clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (TF)n >9 microsatellites in the bovine genome is estimated at >100, COO. The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). The sequence upstream and downstream of the microsatellite sequence were used to generate the required PCR primers for in vitro amplification of the corresp. microsatellite (using the program of PTPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait traits esp. in cattle, to allow selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymorphic bovine DNA markers - used in genetic identification,
                                                                                                                                                                                                                                                                                                                                            PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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0.5%; Score 26.8; DB 13; Length 33;
Best Local Similarity 93.3%; Pred. No. 1.9e+04;
Matches 28; Conservative 0; Mismatches 2; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 33 BP; 0 A; 0 C; 16 G; 17 T; 0 other;
                                                                                                                                                                                                                                                                                                      Microsatellite sequence from clone TGLA415.
                                                                                                                                                                                                                                                                                                                                                                      genetic mapping; traits; amplification; ss.
  2310 CACACACAGACAGACACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           32 CACACACACACACACACACACACACA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene mapping, and selective breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Table 7; Page 334; 517pp; English.
                                                                                                                                                            AAQ34009 standard; DNA; 33 BP.
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                                                                                                                                                                                                                                                         02-FEB-1993 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GENM-) GENMARK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                 Bos taurus.
                                                                                                                                                                                                              AAQ34009;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ34097;
                                                                                                                                       AAQ34009/c
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The sequence is that of a bovine microsatellite sequence obtd.

by screening a library of bovine MboI DNA fragments of between
250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe.
250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe.
250 one out of 50 clones cross-hybridised. Assuming independent
251 distribution of microsatellites and MboI sites, the frequency of
252 (T6) >9 microsatellites in the bovine genome is estimated at >100,
253 ono. The sequence information for ca. 230 such bovine microsatellites
253 summarised in the specification and indexed herein (see below).
254 The sequences upstream and downstream of the microsatellite sequence
255 omericallites may be used to identify individuals,
256 for parentage testing, and in the genetic mapping of economic trait
257 clori, or genes involved the determinism of economically important
258 close, in cattle, to allow selective breeding.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic bovine DNA markers – used in genetic identification, gene mapping, and selective breeding
                                                                                                                                   PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3; bb
1.9e+04;
2;
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                                                                     Sequence of a microsatellite from clone TGLA54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 33 BP; 0 A; 0 C; 16 G; 17 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 92WO-US00340.
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02-FEB-1993 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 93.33 tes 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENM-) GENMARK.
                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                   06-AUG-1992.
                                                                                                                                                                                                                                                                                                 WO9213102-A.
                                                                                                                                                                                                                                       Bos taurus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphism; repeat sequence; genetic marker; primer; amplification; PCR; polymerase chain reaction; paternity; maternity; human; pedigree; linkage analysis; genetic disease; animal; plant; breeding; locus; hybridisation; chromosome; ds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.5%; Score 26.8; DB 18; Length 33; 33.3%; Pred. No. 1.9e+04; Ve 0; Mismatches 2; Indels C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n - using novel nucleic acid mols. as primers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 33 BP; 16 A; 17 C; 0 G; 0 U; 0 other;
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89US-0341562.
94US-0222177.
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94US-0222177.
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nes 28; Conservative
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                                                                                                                                                                                                                                                                                                                                                                         (MARS-) MARSHFIELD CLINIC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-042299/04.
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21-APR-1989;
04-APR-1994;
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                                                                                                                                                                        21-APR-1989;
                                                                                                                                                                                                                                     05-SEP-1991;
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                                   US5582979-A.
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The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n.(dG-dT)n which can be used as genetic markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis. Clones containing the repeat sequences were isolated by hybridisation of probe. Over 100 repeat blocks were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(dC-dA).(dG-dT) probe. Over 100 repeat blocks were isolated. The inserts from the clones were amplified by primers AAT67798-Y66047. Those clones where the repeat sequence has been determined are shown in AAT65704-7797. This repeat sequence is from the marker clone Mdf58 which contains the repeat sequence having the formula: (CA)16.5.
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0.5%; Score 26.8; DB 18; Length 33;
Best Local Similarity 93.3%; Pred. No. 1:9e+04;
Matches 28; Conservative 0; Mismatches 2; Indels (
                                                                                                                                                                                                                                                               Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n - using novel nucleic acid mols. as primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 33 BP; 16 A; 17 C; 0 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Column 11-12; 186pp; English.
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(MARS-) MARSHFIELD CLINIC
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May 31, 2003, 16:41:01; Search time 610 Seconds (without alignments) 11242.271 Million cell updates/sec
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Minimum DB seq length: 10 Maximum DB seq length: 40

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Published_Applications_NA:* Database :

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Sequence 27, Appl	Sequence 678, App	Seguence 93, Appl	Sequence 711, App	Sequence 665, App	Sequence 9, Appli	Sequence 11, Appl	Sequence 1, Appli	Sequence 5, Appli	Sequence 66, Appl	Sequence 770, App	Sequence 600, App	Sequence 11, Appl	Sequence 11, Appl	Sequence 11, Appl	Sequence 21, Appl	Sequence 1, Appli	Sequence 27, Appl	Sequence 3, Appli
SUMMARIES	O.	US-10-085-906-27	US-09-263-959-678	US-10-085-906-93	US-09-263-959-711	US-09-263-959-665	US-09-920-342-9	US-09-920-342-11	US-09-735-363A-1	US-09-735-363A-5	US-09-735-363A-66	US-09-263-959-770	US-09-263-959-600	US-09-970-820-11	us-09-986-718-11	US-09-371-900-11	US-09-931-732-21	US-09-925-674A-1	US-10-231-381-27	US-09-925-674A-3
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Sequence 19, Appl Sequence 19, Appl Sequence 19, Appl Sequence 19, Appl Sequence 506, App Sequence 506, App Sequence 382, App Sequence 393, App Sequence 315, App Sequence 325, App Sequence 325	sequence 1, Appli Sequence 1, Appli Sequence 2, Appli Sequence 173, Ap Sequence 173, Ap Sequence 147, Appl Sequence 147, Appl Sequence 93, Appl Sequence 97, Appl Sequence 97, Appl Sequence 18, Appl Sequence 26, Appl Sequence 26, Appl Sequence 26, Appl Sequence 26, Appl
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Sequence 21, Appl Sequence 155, App Sequence 20, Appl Sequence 1, Appl Sequence 51, Appl Sequence 51, Appl Sequence 51, Appl Sequence 10, Appl Sequence 10, Appl Sequence 10, Appl Sequence 10, Appl Sequence 12, Appl Sequence 10, Appl Sequence 1, Appl Sequence 2, Appl Sequence 2, Appl Sequence 1, Appl	Sequence 50, Appl Sequence 51, Appl Sequence 16, Appl Sequence 20, Appl Sequence 8, Appli Sequence 97, Appl Sequence 19, Appl
9 US-09-835-370-21 9 US-09-888-326-755 9 US-09-888-326-755 9 US-09-9888-326-755 9 US-09-9888-326-756 9 US-09-9888-326-756 9 US-09-9888-326-756 9 US-09-988-326-756 9 US-09-988-326-759 9 US-09-988-326-759 9 US-09-988-326-759 9 US-10-017-995-54 9 US-10-017-995-54 9 US-10-017-995-54 9 US-10-017-995-54 9 US-10-017-995-54 9 US-10-017-995-54 9 US-09-776-479-91 9 US-09-965-116A-79 9 US-09-776-479-91 9 US-09-965-116A-79 9 US-09-75A-75A-75A-75 9 US-09-75A-75A-75 9 US-09-75A-75A-75 9 US-09-75A-75A-75 9 US-09-75A-75A-75 9 US-09-75A-75A-75 9 US-09-75A-75A-75	US-09-500-700-21 US-09-500-700-21 US-09-934-060A-16 US-09-934-060A-16 US-09-744-41-20 US-10-135-965-8 US-10-135-965-51 US-10-135-965-51 US-10-035-041-19 US-10-035-041-19 US-10-035-19-19-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-119-19 US-10-035-
US-09-835-1770-21 US-09-888-136-755 US-09-888-136-755 US-09-888-136-755 US-09-818-138-138-5 US-09-818-138-5 US-09-818-138-5 US-09-818-138-5 US-10-112-653-85 Sequence US-10-112-653-85 Sequence US-10-112-653-85 Sequence US-10-112-653-85 Sequence US-10-112-653-85 Sequence US-10-112-653-85 Sequence US-09-776-479-13 Sequence US-09-776-479-91 Sequence US-09-814-468-104 US-09-814-468-104 US-09-814-468-104 US-09-814-468-104 US-09-965-116A-79 Sequence US-09-965-116A-99 Sequence US-09-	9 US-09-500-705-125 Sequence 9 US-09-848-754A-5701 Sequence 10 US-09-714-414-20 Sequence 9 US-10-135-965-8 Sequence 10 US-10-135-965-51 Sequence 10 US-10-135-965-51 Sequence 10 US-10-036-041-19 Sequence 10 US-10-036-214-19 Sequence 10 US-10-036-160-19 Sequence 10 US-10-036-160-19 Sequence
9 US-09-835-370-21 Sequency 9 US-09-888-326-755 Sequency 9 US-09-9888-326-755 Sequency 9 US-09-931-732-20 Sequency 9 US-09-931-732-20 Sequency 9 US-09-931-732-20 Sequency 9 US-10-112-653-85 Sequency 9 US-10-112-653-85 Sequency 9 US-10-017-995-54 Sequency 9 US-10-017-995-54 Sequency 9 US-10-017-995-55 Sequency 9 US-10-017-995-55 Sequency 9 US-10-017-995-55 Sequency 9 US-09-776-479-91 Sequency 9 US-09-965-116A-98 Sequency 9 US-09-965-116A-98 Sequency 9 US-09-965-116A-99 Sequency 9 US-09-965-10A-90 Sequency	3 8 9 05-09-500-700-51 Sequence 3 18 9 05-09-848-7548-5701 Sequence 3 18 9 05-09-934 Co6A-16 Sequence 3 18 9 05-09-934 Co6A-16 Sequence 3 18 9 05-09-934 Co6A-16 Sequence 3 18 0 05-09-774-414-20 Sequence 3 18 0 05-10-135-965-37 Sequence 3 18 0 05-10-135-965-31 Sequence 3 18 0 05-10-035-855-19 Sequence 3 18 0 05-10-035-855-19 Sequence 3 18 0 05-10-035-119 Sequence 3 18 0 05-10-035-119 Sequence 3 18 0 05-10-036-160-19 Sequence 3 0 05-10-036-160-19 Sequence 3 0 05-10-036-160-19 Sequence 3 0 05-10-036-160-19 Sequence 3 0 05-10-
18         0.4         18         9         US-09-883-376-755         Sequence           18         0.4         18         9         US-09-883-376-755         Sequence           18         0.4         18         9         US-09-931-732-20         Sequence           18         0.4         18         9         US-09-818-936-755         Sequence           18         0.4         18         9         US-10-112-653-1         Sequence           18         0.4         18         9         US-10-112-653-1         Sequence           18         0.4         18         9         US-10-117-955-1         Sequence           18         0.4         18         9         US-10-117-955-5         Sequence           18         0.4         18         9         US-10-107-955-5         Sequence           18         0.4         18         9         US-10-107-95-5         Sequence	8 0.3 38 9 05-09-500-700-51 Sequence 8 0.3 38 9 05-09-500-700-51 Sequence 8 0.3 38 9 05-09-934 060A-16 Sequence 8 0.3 39 10 05-09-934 060A-16 Sequence 8 0.3 40 9 05-10-135-965-8 Sequence 8 0.3 40 9 05-10-135-965-37 Sequence 10 0.3 24 9 05-10-135-965-19 Sequence 10 0.3 24 9 05-10-036-041-19 Sequence 10 0.3 24 9 05-10-036-24-19 Sequence 10 0.3 24 9 05-10-036-24-19 Sequence 10 0.3 24 9 05-10-036-214-19 Sequence 10 0.3 24 9 05-10-036-214-19 Sequence 10 0.3 24 9 05-10-036-214-19 Sequence 11 0.3 24 9 05-10-036-214-19 Sequence 12 0.3 24 9 05-10-036-214-19 Sequence 13 0.3 24 9 05-10-036-214-19 Sequence 14 0.3 24 9 05-10-036-214-19 Sequence 15 0.3 24 9 05-10-036-214-19 Sequence 16 0.3 24 9 05-10-036-160-19 Sequence

Sequence 16, Appl Sequence 1525, App Sequence 1624, App Sequence 1624, App Sequence 2017, App Sequence 2017, App Sequence 2017, Appl Sequence 2017, Appl Sequence 20, Appl Sequence 20, Appl Sequence 20, Appl Sequence 215, Appl Sequence 217, Appl Sequence 218, Appl Sequence 218, Appl Sequence 218, Appl Sequence 218, Appl Sequence 219, Appl Sequence 229, Appl Sequence 220, Appl Sequence 230, Appl Sequence 230, Appl Sequence 230, Appl Sequence 230, A	
36 10 US-09-306-417-16 37 9 US-09-908-504A-37 38 9 US-09-930-423-1868 38 9 US-09-930-423-1868 38 9 US-09-930-423-1868 38 9 US-09-930-423-2017 38 9 US-09-930-423-2017 39 9 US-10-067-956-40 39 9 US-10-067-956-40 39 10 US-09-933-452-22 39 10 US-09-915-060-48 40 10 US-09-915-061-33 50 10 US-09-18-08-32 50 10 US-09-18-08-32 50 10 US-09-18-08-33 50 10 US-09-734-847-45 50 10 US-09-734-847-65 50 10 US-09-734-847-65 50 10 US-09-734-847-65 50 10 US-09-734-847-67 50 10 US-09-912-263-3 50 10 US-09-912-263-3 51 10 US-	
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Sequence 191, App Sequence 3881, Ap Sequence 27, Appl Sequence 17, Appl Sequence 17, Appl Sequence 17, Appl Sequence 17, Appl Sequence 11, Appl Sequence 11, Appl Sequence 19, Appl Sequence 19, Appl Sequence 10, Appl Sequence 24, Appl Sequence 24, Appl Sequence 24, Appl Sequence 24, Appl Sequence 29, Appl Sequence 29, Appl Sequence 29, Appl Sequence 29, Appl Sequence 21,	sequence 3670, Sequence 1555, Sequence 1, M. Sequence 18, A. Sequence 18, A. Sequence 18, A. Sequence 13, Sequence 105, Sequence 124, Sequence 124, Sequence 14, Sequence 14, Sequence 4, M. Sequence 5, M. Sequence 6,
9 US-09-992-665-191 25 10 US-09-866-108-3881 25 10 US-09-866-108-3881 26 US-09-866-108-3881 27 US-09-866-108-3881 28 US-09-33-052-37 30 US-09-33-527-12 30 US-09-33-527-12 31 US-09-992-220B-89 31 US-09-999-220B-95 31 US-09-999-220B-95 31 US-09-999-220B-95 31 US-09-999-220B-95 31 US-09-999-220B-95 31 US-09-999-220B-95 31 US-09-999-220B-95 32 US-09-999-220B-95 33 US-09-999-220B-95 34 US-09-994-94-10 36 US-09-944-91-10 36 US-09-944-852-10 36 US-09-944-857-10 36 US-09-944-857-10 36 US-09-944-857-10 36 US-09-944-877-10 36 US-09-944-877-10 37 US-09-944-877-10 38 US-09-943-811A-10 39 US-09-955-372-374 40 US-09-958-372-374 40 US-09-96-277-29 40 US-09-98-838-1755 25 US-09-730-289B-1755 25 US-09-730-289B-1755	25 9 US-09-730-289B-3670 25 10 US-09-730-289B-5670 26 10 US-09-730-881-1 27 9 US-09-86-18-1555 27 9 US-09-87-76-881-1 27 10 US-09-84-23 30 9 US-09-984-848 31 10 US-09-981-848 31 10 US-09-981-15-105 31 10 US-09-91-263-105 31 10 US-09-91-274-1561 33 10 US-09-801-274-1561 33 10 US-09-801-274-1561 34 10 US-09-862-847-4
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Sequence 3480, Ap Sequence 3629, Ap Sequence 3851, Ap Sequence 3908, Ap Sequence 1541, Ap Sequence 1186, Ap Sequence 1575, Ap Sequence 42, Appl Sequence 43, Appl Sequence 43, Appl	Sequence 15, Appl Sequence 16, Appl Sequence 17, Appl Sequence 10, Appl Sequence 15, Appl Sequence 2005, Appl Sequence 4, Appl Sequence 4, Appl Sequence 17, Appl Sequence 4, Appl Sequence 17, Appl Sequence 11, Appl Sequence 12, Appl Sequence 12, Appl Sequence 12, Appl Sequence 12, Appl Sequence 13, Appl Sequence 12, Appl Sequence 13, Appl Sequence 13, Appl Sequence 14, Appl Sequence 11, Appl Sequence 11, Appl Sequence 12, Appl Sequence 13, Appl Sequence 14, Appl Sequence 14, Appl Sequence 15, Appl Sequence 17, Appl Sequence 21, Appl Sequence 22, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 22, Appl Sequence 23, Appl Appl Sequence 23, Appl Appl Sequence 23, Appl Appl Sequence 23, Appl Appl S
US-09-780-533A-3480 US-09-780-533A-3629 US-09-780-533A-3621 US-09-780-533A-3908 US-09-780-164-1186 US-09-780-164-1186 US-09-1818-879-42 US-09-8118-879-42 US-09-8118-879-43 US-09-811-7558-42 US-09-211-7558-43 US-09-211-7558-43	10 US-09-888-049-4  11 US-09-963-667-15  12 US-09-963-667-16  13 US-09-963-667-18  14 US-10-135-965-61  15 US-10-135-965-61  16 US-09-998-425-17  17 US-10-135-965-61  18 US-10-137-788-16  19 US-09-997-81-10  10 US-09-967-718-10  10 US-09-967-718-10  11 US-10-135-965-61  12 US-10-137-788-16  13 US-10-137-788-16  14 US-10-137-788-16  15 US-10-137-788-16  16 US-09-730-2898-1754  17 US-09-730-2898-1754  18 US-09-730-2898-1754  19 US-09-730-2898-1754  10 US-09-730-2898-1754  10 US-09-730-2898-1754  10 US-09-86-108-3885  10 US-09-730-2898-1668  10 US-09-730-2898-1754  10 US-09-86-108-18  10 US-09-194-13  10 US-09-191-31  10 US-09-191
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Sequence 711, Application US/09263959
Patent No. US20020150891A1
GENERAL INFORMATION:
APPLICANT: Rowen, Lee
APPLICANT: Rowen, Lee
APPLICANT: Roy, Ban F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH INUMBER OF SEQUENCES: 1279
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                              0.5%; Score 25.8; DB 9; Length 30; 93.1%; Pred. No. 5.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Seed and Berry LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
CITY: Seattle
STATE: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.5%; Score 25.2; DB 10; 90.0%; Pred. No. 8.3e+03;
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                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
CURRENT FILING DATE: 2002-02-27
PRIOR APPLICATION NUMBER: US 60/126,215
PRIOR FILING DATE: 1999-03-25
PRIOR FILING DATE: 1999-03-24
PRIOR FILING DATE: 2000-03-24
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 545
SOFTWARE: FASTSEQ for Windows Version 4.0
SED ID NO 93
LENGTH: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: McMasters, David D.
REGISTRATION UNBER: 33,963
REFERENCE/DOCKET UNMER: 920010.426C2
TELECOMMUNICATION INFORMATION:
TELEPAX: (206) 62-4900
TELEFAX: (206) 68-2631
INFORMATION FOR SEQ ID NO: 711:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
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                                                                                                                                                                                                                                                                                                            Query Match 0.5%
Best Local Similarity 93.1%
Matches 27; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                 TYPE: DNA
CORGANISM: Homo sapiens
US-10-085-906-93
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 4
US-09-263-959-711/c
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US-09-263-959-711
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                                                                                                                                                                                                                                                                                                                                                                                                                                       qq
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APPLICANT: Hood, Lercy E.
APPLICANT: Rowen, Lee
APPLICANT: Rowen, Lee
APPLICANT: Roop, Ben F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
NUMBER OF SEQUENCES: 1279
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Publication No. US20030054371A1
GENERAL INFORMATION:
APPLICANT: Ying, Vincent
APPLICANT: Wu, Paul
APPLICANT: Gray, Gary, Sary
TITLE OF INVENTION: POLYMORPHIC ELEMENTS IN THE
TITLE OF INVENTION: COSTIMULATORY RECEPTOR LOCUS AND USES THEREOF
CURRENT APPLICATION NUMBER: US/10/085,906
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                                                 3; DB 9; Length 31; 3e+03;
                                                                                         2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: US/MR-1999
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Seed and Berry LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 26.8; DB 10;
Pred. No. 3.5e+03;
                                                 0.5%; Score 26.8; C
93.3%; Pred. No. 3e+0
tive 0; Mismatches
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                                                                                                                                 2310 CACACACACAGACAGACACACACACACA 2339
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                                                                                                                                                         31 CACACACACACACACACACACACACA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                     US-09-263-959-678/c; Sequence 678, Application US/09263959; Patent No. US20020150891A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: MCMASLETS, David D.
REGISTRATION NUMBER: 33,963
REFRENCE/POCKET NUMBER: 92001
TELECOMMUNICATION INFORMATION:
TELEFAX: (206) 682-693
INFORMATION FOR SEQ ID NO: 678:
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LENGTH: 39 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 93.35
Matches 28; Conservative
                                                                      Local Similarity 93.3 nes 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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US-09-263-959-678
            US-10-085-906-27
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                                                     Query Match
                                                                                           Matches
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APPLICANT: Filion, Mario
APPLICANT: Filion, Mario
APPLICANT: Philip, Nigel
TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides
TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides
FILE REFERENCE: 02811-0181
CURRENT APPLICATION NUMBER: US/09/735,363A
CURRENT FILING DATE: 2000-12-12
PRIOR FILING DATE: 1999-12-13
PRIOR FILING DATE: 1999-12-13
PRIOR PLICATION NUMBER: 60/228,925
PRIOR PLILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 87
SOFTWARE: Patentin version 3.0
SEQ ID NO 1
LENGTH: 27
                                                                                                                                Gaps
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Fatent No. US/20020137709A1
Fatent No. US/20020137709A1
Fatent No. US/20020137709A1
FAPLICANT: University of Southern California
APPLICANT: Lin, Shi-Lung
APPLICANT: Chuong, Cheng-Ming
APPLICANT: Chuong, Cheng-Ming
APPLICANT: Chuong, Cheng-Ming
APPLICANT: Chuong, Cheng-Ming
APPLICANT: University of Southern California
FILE REFERENCE: 13761-7024
CURRENT APPLICATION UNMER: US/09/920, 342
CURRENT FILING DATE: 2002-01-17
PRIOR PAPLICATION NUMBER: US 60/222, 479
PRIOR PAPLICATION NUMBER: US 60/222, 479
PRIOR PAPLICATION NUMBER: US 60/222, 479
FRIOR FILING DATE: 2000-08-02
SOFTWARE FRESEQ for Windows Version 4.0
SEQ ID NO 11
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                                                                                                                          0; Indels
                                                                      0.5%; Score 24; DB 10; I 100.0%; Pred. No. 1.3e+04; tive 0; Mismatches 0;
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Pred. No. 1.6e+04;
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Best Local Similarity 100.0%; Pred. No. 1.3e+04;
Matches 24; Conservative 0; Mismatches 0;
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; OTHER INFORMATION: Anti-bcl2 primer US-09-920-342-9
                                                                                                                                                                          2725 CCATGCCTCCCTGGCCTGAAGAAG 2748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1985 GGATGACTGAGTACCTGAACCGGC 2008
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                                                                                                                                                                                                      24 CCAIGCCICCCIGGCCIGAAGAAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: bc12 primer US-09-920-342-11
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                                                                         Query Match 0.5%
Best Local Similarity 100.0
Matches 24; Conservative
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Best Local Similarity
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US-09-735-363A-1/C
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US-09-920-342-11
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                                                                 Sequence 665, Application US/09263959
Patent No. US/20020150891A1
GENERAL INFORMATION:
APPLICANT: ROWEN, Lee
APPLICANT: ROWEN, Lee
APPLICANT: ROOP, Ben F
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI
NUMBER OF SEQUENCES:
ADDRESSEE: Seed and Berry LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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APPLICANT: Lin, Shi-Lung
APPLICANT: Chuong, Cheng-wing
APPLICANT: Chuong, Cheng-wing
TITLE OF INVENTION: Randahla.
FILE REFERENCE: 13761-7024
CURRENT APPLICATION NUMBER: US/09/920,342
CURRENT APPLICATION NUMBER: US 60/222,479
PRIOR FILING DATE: 2000-08-02
NUMBER OF SEQ ID NOS: 15
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 24
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MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPOTER: IBM PC COMPATIBLE
COMPOTER: IBM PC COMPATIBLE
COMPOTER: A PATENTIN Release #1.0, Version #1.25
SOFTWARE PATENTION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE:
                                                                                                                                                                                                                                                                                               E: Seed and Berry LLP
6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.5%; Score 24.2; DB 10;
Best Local Similarity 89.7%; Pred. No. 1.3e+04;
Matches 26; Conservative 0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: McMasters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.426C2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2310 CACACACAGACAGACACACACACACAC 2338
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Patent No. US20020137709A1
GENERAL-INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 665:
SEQUENCE CHARACTERISTICS:
LENGTH: 29 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
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CITY: Seattle
STATE: Washington
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                                         US-09-263-959-665/c
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US-09-263-959-665
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Seattle
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                                               US-09-263-959-770/c
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                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                          STREET:
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                                                                                                                                                                                    | Sequence 5, Application US/09735363A | Sequence 5, Application US/09735363A | Sequence 5, Application US/0010041681A1 | Settent No. US20010041681A1 | Seneral Information: | Application | Marion | Application | Marion | Application | Marion | TILLE REFERENCE: 02811-011 | FILLE REFERENCE: 02811-011 | FILLE REFERENCE: 02811-011 | FILLE REFERENCE: 02811-011 | Settent Application Number: 05/0170,325 | PRIOR FILLING DATE: 1999-12-13 | PRIOR FILLING DATE: 1999-12-13 | PRIOR FILLING DATE: 2000-08-29 | NUMBER OF SEQ ID NOS: 87 | SettentIn Version 3:0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: FILLOW, Wigel
APPLICANT: FILLOW, Wigel
TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides
TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides
FILE REFERENCE: 02811-0181
CURRENT APPLICATION NUMBER: 60/9735,363A
CURRENT FILING DATE: 2000-12-12
PRIOR FILING DATE: 1999-12-13
PRIOR FILING DATE: 1999-12-13
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 87
SOFTWARE: PATENTIN VERSION 3.0
Gaps
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Best Local Similarity 92.6%; Pred. No. 1.6e+04;
Matches 25; Conservative 0; Mismatches 2; Indels
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0.5%; Score 23.8; DB 10; Length 27;
Best Local Similarity 92.6%; Pred. No. 1.6e+04;
Matches 25; Conservative 0; Mismatches 2; Indels 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: Synthetic Oligonucleotide US-09-735-363A-5
       0; Mismatches
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                                               2310 CACACACACAGACAGACACACACAC 2336
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APPLICANT: Fillon, Mario
                                                                         27 CACACACACACACACACACACACAC 1
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           25; Conservative
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                   Matches
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Sequence 770, Application US/09263959
Sequence 770, Application US/09263959
Patent No. US20020150891A1
GENERAL INFORMATION:
APPLICANT: HOWO, Leroy E.
APPLICANT: ROOP, Ben F.
TITIE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UNTHERE OF ESCUENCES: 1279
CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed and Berry LLP
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Patent No. US20020150891A1

GENERAL INFORMATION:

APPLICANT: Hood, Leroy E.

APPLICANT: Rowen, Lee

CORRESCOUDENCES: 1279

CORRESCONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.5%; Score 23.8; DB 10; 92.6%; Pred. No. 1.6e+04;
                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: TIM PC Compatible
COMPUTER: TIM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
APPLICATION DATA: US/09/263,959
FILING DATE: ..05-MAR-1999
                                                                                                                                                                                                                                 5: Seed and Berry LLP
6300 Columbia Center, 701 Fifth Avenue
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: MCMASTERS, DAVID D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (206) 622-4900
TELEPAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 770:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 92.6 Matches 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 27 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98104-7092
                                                                                                                                                                                                                                                                                                               STATE: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seattle
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
APPLICATION DATA:
FILING DATE: 05-OCE-2001
PRIOR APPLICATION: <u cutoffcent con control con
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSE: PENNIE & EDMONDS
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/386,844
FILING DATE: 10-FEB-1995
ATTORNEY,AGENT INFORMATION:
NAME: COLUZZI, LAULA A.
REGISTRATION NUMBER: 30,742
REFERRUCE/DOCKET NUMBER: 7853-032
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPAS: (212) 869-8864
                                                                                                                                                                                                                                                    33,963
33,963
mmp. 920010.426C2
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                                                                                 MBER: US/09/263,959
05-MAR-1999
                                                                                                                                                                      ATTORNEY AGENT INFORMATION:
NAME: MCMASters, David D.
REGISTRATION NUMBER: 33,963
REFRENCE/DOCKET NUMBER: 92001
TELECOMMUNICATION INFORMATION:
TELEFAN: (206) 622-4900
TELEFAN: (206) 682-6931
INFORMATION FOR SEQ ID NO: 600:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence II, Application US/09970820
Patent No. US20020170077A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: FALB, DEAN A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New York
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                                                                                                                                                  CLASSIFICATION:
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                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY:
US-09-263-959-600
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WL INFORMATION.
APPLICANT: FALB, DEAN A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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                                                                                                                                                  Length 24;
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ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUN TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: DATE: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
APPLICATION NUMBER: US/09/986,718
FILING DATE: 09-No. US20020178458A1-2001
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION 
                                                                                                                                                                                                                        Indels
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95.8%; Pred. No. 3.4e+04;
tive 0; Mismatches 1; Indels
                                                                                                                  Query Match
0.4%; Score 22.4; DB 9;
Best Local Similarity 95.8%; Pred. No. 3.4e+04;
Matches 23; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 1155 Avenue of the Americas CITY: New York STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 7853-032
TELECOMAUNICATION INCORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: PENNIE & EDMONDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION
NAME: CORUZZI, LAURA A.
REGISTRATION NUMBER: 30,742
                                                                                                                                                                                                                                                                1875 GGACGGGTGAACTGGGGGAGGAT 1898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 11, Application US/09986718; Patent No. US20020178458A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                     1875 GGACGGGTGAACTGGGGGAGGAT 1898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 11, Application US/09371900; Patent No. US20020137700A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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hes 23; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 14
US-09-986-718-11/c
                                US-09-970-820-11
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US-09-371-900-11/c
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1884 GAACTGGGGGAGGATTGTGGCCTTCTT 1910
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: 16
OTHER INFORMATION: n is inosine
NAME/KEX: modified_base
LOCATION: 19
OTHER INFORMATION: n is inosine
NAME/KEY: modified_base
LOCATION: 22
                                                                                                                                                                                                                                                                1722 GGTGCCACCTGTGGTCCACCTG 1743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: n is inosine NAME/KEY: modified_base LOCATION: 25
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                                                                                                                                                                                                                                                                                          1 GGTGCCACCTGTGGTCCACCTG 22
                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Mouse
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US-10-231-381-27
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LENGIH: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA
                                  SEQ ID NO 21
                                                                              TYPE: DNA
                                                            LENGIH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           δλ
                                             TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT AND DIAGNOSIS OF CARDIOVASCULAR DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: BIOWN, BOD D.
APPLICANT: BIOWN, BOD D.
APPLICANT: RILBY, TIMOTHY A.
APPLICANT: RILBY, TIMOTHY A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES COMPRISING
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES COMPRISING
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES COMPRISING
TITLE OF INVENTION: ANTISENSE DASS
TITLE OF INVENTION: UNUMBER: US/09/931,732
CURRENT APPLICATION NUMBER: PCT/US00/09293
PRIOR PILING DATE: 2000-04-07
PRIOR FILING DATE: 2000-04-07
PRIOR FILING DATE: 1999-04-08
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0.4%; Score 22.4; DB 10; Length 24;
Best Local Similarity 95.8%; Pred. No. 3.4e+04;
Matches 23; Conservative 0; Mismatches 1; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DESCRIPTION: /desc = "synthetic oligonuclectide"
                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/599,654
FILING DATE: 09-FEB-1996
APPLICATION NUMBER: US 08/485,573
FILING DATE: 07-JUN 11995
APPLICATION NUMBER: US 08/386,844
APPLICATION NUMBER: US 08/386,844
FILING DATE: 10-FEB-1995
ATORNEY/AGENT INFORMATION:
NUMBER: CORUZZI, LAURA A
REGISTRATION NUMBER: 30,742
REFERENCE/COCKET WUMBER: 7853-104
                                                                                                                                                            STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE DESCRIPTION: SEQ ID NO: 11: US-09-371-900-11
                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: PENNIE & EDMONDS LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864
TELEFAX: 6141 PENITE
INFORMATION FOR SEQ 1D NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1875 GGACGGGGTGAACTGGGGGAGGAT 1898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 16
US-09-931-732-21
Sequence 21, Application US/09931732
Publication No. US20030045488A1
GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                          NUMBER OF SEQUENCES: 54
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                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 10036-2711
                                                                                                                                                                                                              STATE: New York
                                                                                                                                                                                               New York
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Sequence 1, Application US/09925674A

Patent No. US20020119943A1

GENERAL INFORMATION:
TITLE OF INVENTION: A MOVEL MAMMALIAN GENE, bcl-w, BELONGS TO THE bcl-2

TITLE OF INVENTION: A MOVEL MAMMALIAN GENE, bcl-w, BELONG TO THE bcl-2

TITLE OF INVENTION: A MOVEL MAMMALIAN OF APOPTOSIS-CONTROLLING GENES

TITLE OF INVENTION: PAMILY OF APOPTOSIS-CONTROLLING GENES

FILE REFERENCE: 11666a

CURRENT FILING DATE: 2001-08-09

PRIOR APPLICATION NUMBER: 09/925,674

PRIOR PILING DATE: 2001-08-09

PRIOR PILING DATE: 1996-03-27

NUMBER: OF THE NUMBER: PN8965

PRIOR ELING DATE: 1996-03-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
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                                                                                                                                                                 ö
                                                                                              Query Match 0.4%; Score 22; DB 9; Length 22; Best Local Similarity 100.0%; Pred. No. 4e+04; Matches 22; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Chen, Xiangning TITLE OF FOR GENOTYPING TITLE OF INVENTION: ALLELE SPECIFIC PCR FOR GENOTYPING FILE REFERENCE: 02940201aa CURRENT APPLICATION NUMBER: US/10/231,381 CURRENT FILING DATE: 2002-08-30
OTHER INFORMATION: Synthetic oligonucleotide primers US-09-931-732-21
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GENERAL INFORMATION

APPLICANT: AMRAD OPERATIONS PLY Ltd

TITLE OF INVENTION: A NOVEL MAMMALIAN GENE, bcl-w, BELONGS TO THE bcl-2

TITLE OF INVENTION: FAMILY OF APOPTOSIS-CONTROLLING GENES

FILE REFERENCE: 11686a

CURRENT APPLICATION NUMBER: US/09/925,674A

PRIOR FILING DATE: 2001-08-09

PRIOR FILING DATE: 2001-08-09

PRIOR PLICATION NUMBER: PN8965

NUMBER OF SEQ ID NOS: 9

SOFTWARE: Patentin Ver: 2.1

**ENAME: 11
                                                                                                                                                                                                                                                                                                  Gaps
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0.4%; Score 21.2; DB 9; Length 39;
Best Local Similarity 76.5%; Pred. No. 9.6e+04;
Matches 26; Conservative 0; Mismatches 8; Indels
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                                                                                                                                                                         ; OTHER INFORMATION: Synthetic oligonucleotide primer US-10-231-381-27
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80.8%; Pred. No. 9.2e+04;
tive 1; Mismatches 4;
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APPLICANT: Sloley, David R.
APPLICANT: Hombin, Mark
TITLE OF INVENTION: The ST-B17 Serotonin Receptor
FILE REFERENCE: NIH047.1CPIC1
                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/315,776
PRIOR FILING DATE: 2001-08-30
NUMBER OF SEQ ID NOS: 44
SCOTTAMRE: Patentin version 3.1
LENGTH: 39
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LOCATION: 14
COTHER INFORMATION: n is inosine
LOCATION: 17
COTHER INFORMATION: n is inosine
LOCATION: 17
COTHER INFORMATION: n is inosine
LOCATION: 20
COTHER INFORMATION: n is inosine
US-09-925-674A-3
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; Sequence 3, Application US/09829631A
; Patent No. US20020091233A1
                                                                                                                TYPE: DNA
ORGANISM: Artificial sequence
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WS-09-925-674A-3/C
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APPLICANT: Krieg, Daniel J.
TITLE OF INVENTION: IMMUNOSTIMULATORY NUCLEIC ACID FOR
TITLE OF INVENTION: TREATMENT OF NON-ALLERGIC INFLAMMATORY DISEASES
FILE REFERENCE: CO1039/70060(AMS)
CURRENT APPLICATION NUMBER: US/10/112,653
PRIOR APPLICATION NUMBER: US 60/279,642
                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 21
US-09-764-891-10175/c

Sequence 10175, Application US/09764891

Sequence 10175, Application US/09764891

GENERAL INFORMATION:

APPLICANT: ROSEN et al.

TITLE OF INVENTION: Wacleic Acids, Proteins, and Antibodies

CURRENT APPLICATION NUMBER: US/09/764,891

FILOR APPLICATION NUMBER: US/09/764,891

FILOR APPLICATION ADATE: 2001-01-17

NUMBER OF SEQ ID NOS: 10231

SEQ ID NO 10175

LENGTH: 38
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82.8%; Pred. No. 9.6e+04;
tive 0; Mismatches 5; Indels
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73.0%; Pred. No. 1.1e+05;
tive 0; Mismatches 10; Indels
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CURRENT APPLICATION NUMBER: US/09/829,631A CURRENT FILING DATE: 2001-04-10 PRIOR APPLICATION NUMBER: US 08/428,242 PRIOR FILING DATE: 1995-09-18
                                                                           NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Synthetic Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                             31 GGCTGGCCTTCAAGGCCTTCCTGCTATGC 3
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1012
LENGTH: 24
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Publication No. US20030050268A1
GENERAL INFORMATION:
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OTHER INFORMATION: primer
US-09-829-631A-3
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ORGANISM: Artificial Sequence
FEATURE:
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Best Local Similarity 73.09
Matches 27; Conservative
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US-09-764-891-10175
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US-10-112-653-1012/c
                                                                                                                    SEQ ID NO 3
LENGTH: 33
                                                                                                                                                              TYPE: DNA
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Matches
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Sequence 22, Application US/09735363A

Sequence 22, Application US/09735363A

Patent No. US20010041681A1

Patent No. US20010041681A1

GENERAL INFORMATION

APPLICANT: Fillin, Mario

APPLICANT: Fillin, Migel

APPLICANT: Fillin, Migel

APPLICANT: Phillip, Nigel

APPLICANT: Phillip, Nigel

TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides

TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides

CURRENT APPLICATION NUMBER: US/09/735,363A

CURRENT FILING DATE: 1099-12-13

PRIOR FILING DATE: 1999-12-13

PRIOR APPLICATION NUMBER: 60/728,925

PRIOR PELICATION NUMBER: 60/228,925

PRIOR FILING DATE: 2000-08-29

NUMBER OF SEQ ID NOS
                                                                                            APPLICANT: Filion, Mario
APPLICANT: Filion, Mario
APPLICANT: Filion, Migel
APPLICANT: Filion, Migel
APPLICANT: Filion, Migel
TITLE OF INVENTION: Therapeutically Useful Synthetic Oligonucleotides
TITLE REFERENCE: 02811-0181
FILE REFERENCE: 02811-0181
CURRENT APPLICATION NUMBER: 06/170,325
PRIOR APPLICATION NUMBER: 60/170,325
PRIOR FILING DATE: 1999-12-13
PRIOR FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 87
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.4%; Score 20.8; DB 10; Length 24; Best Local Similarity 91.7%; Pred. No. 8.6e+04; Matches 22; Conservative 0; Mismatches 2; Indels (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Synthetic Oligonucleotide US-09-735-363A-22
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US-09-735-363A-21/c
US-09-735-363A-21/c
Sequence 21, Application US/09735363A
Setent No. US20010041681A1
GENERAL INFORMATION:
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US-10-085-906-189/C
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; Sequence 1068, Application US/09776479

; Dublication No. US20030087848A1

; Publicant Bratzler, Robert L.

APPLICANT: Petersen, Deanna M.
APPLICANT: Petersen, Deanna M.
APPLICANT: Fouron, Yves

TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the
TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the
TITLE OF INVENTION: Treatment of Acthma and Allergy

FILE REFERENCE: C1037/7013 (HCL/MAT)

FULE REPERENCE: 2001-02-02

CURRENT FILING DATE: 2001-02-02

; CURRENT FILING DATE: 2001-02-03

; PRIOR FILING DATE: 2000-02-03

; NUMBER OF SED ID NOS: 1093

; SEQ ID NO 1068

; SEQ ID NO 1068
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                 0.4%; Score 20.8; DB 9; Length 24; 91.7%; Pred. No. 8.6e+04; tive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                         NESTLO 17-995-1068/c NESTLO 17-995-1068/c Sequence 1068, Application US/10017995 Publication No. US20030055014Al GENERAL INFORMATION:
                                                                                                                        2310 CACACACACAGACAGACACACA 2333
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                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                  Query Match
Best Local Similarity 91.7%
Matches 22; Conservative
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Best Local Similarity
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US-09-776-479-1068/c
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US-10-112-653-1012
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Gaps

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Gaps

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Indels

Length 24;

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Gaps
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O.4%; Score 20.8; DB 9; Length 34;
Best Local Similarity 78.1%; Pred. No. 1.1e+05;
Matches 25; Conservative 0; Mismatches 7; Indels
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APPLICANT: Brown, Bob D.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES COMPRISING
FILE REFERENCE: OASBIO.00101
CURRENT APPLICATION UNUBERS US/09/931,732
PRIOR APPLICATION NUMBER: US/09/931,732
PRIOR APPLICATION NUMBER: PCT/USO/09293
PRIOR PILING DATE: 2001-08-16
PRIOR PLILING DATE: 1999-04-08
SOFTWARE: PASCED FOR THIRD OF SEQ ID NOS: 30
SOFTWARE: PASCED FOR WINDOWS VETSION 4.0
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                                                                                                                                                                                                                                                                                                                                                                                     7; Indels
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                                                                                                                                                                                                                                                                                                                                                                                              2312 CACACACAGACAGACACACACACAGAA 2343
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 189, Application US/10085906
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Sequence 22, Application US/09931732
Publication No. US20030045488A1
GENERAL INFORMATION:
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           Publication No. US20030054371A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Best Local Similarity 95.5%
Matches 21; Conservative
                                                                                                                                                                                                                                                                                   ;) TYPE: DNA
ORGANISM: Homo sapiens
US-10-085-906-189
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APPLICANT: Lin, Shi-Lung
APPLICANT: Lin, Shi-Lung
APPLICANT: Lin, Shi-Lung
APPLICANT: Ying, Shao-Yao
TITLE OF INVENTION: Method for Thermocycling Amplification of Nucleic Acid Seque
TITLE OF INVENTION: Generation of Related Peptides Thereof
TITLE OF INVENTION: Generation of Related Peptides Thereof
CURRENT APPLICATION NUMBER: US/09/949,305B
PRIOR APPLICATION NUMBER: 09/494,212
PRIOR APPLICATION NUMBER: 09/494,212
NUMBER OF SEQ ID NOS: 12
SOFWARE: Patentin version 3.1
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1S-69-25
1S-69-25
1S-69-25
1S-69-25
1Publication No. US20030073085a1
1Publication No. Dalxing
1Publication No. Amplifying Expressed Sequences From Genomic DNA OF HIGHER-ORDE
1PILE OF INVENTION: ENGRARYORIC ORGANISMS FOR DNA ARRAYS
1PILE OF INVENTION: US201-290
1PILE OF INVENTION: US2
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GENERAL INFORMATION:
APPLICANT: Max-Delbruck-Centrum fur Molekulare Medizin
TITLE OF INVENTOR: Chimeric Oligonucleotides and the Use Thereof
CURRENT FILING DATE: 101195-24
CURRENT FILING DATE: 2001-03-26
PRIOR APPLICATION NUMBER: DE 197 20 151.2
NUMBER: PRIOR APPLICATION NUMBER: DE 197 20 151.2
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: antisense bcl-2 primer US-09-949-305B-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11, Application US/09817387
Patent No. US20010039263A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2729 GCCTCCCTGGCCTGAAGAAG 2748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: artificial sequence
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-817-387-11/c
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APPLICANT: Hood, Leroy E.
APPLICANT: Rowen, Lee
APPLICANT: Koop, Ben F.
TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UNDRER OF SEQUENCES: 1279
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                       Gaps
                          ; OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid US-09-864-785-1826
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                                                                                                       Query Match 0.4%; Score 20; DB 9; Length 37; Best Local Similarity 69.4%; Pred. No. 1.9e+05; Matches 25; Conservative 1; Mismatches 10; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
VITORNEY AAAR.
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91.3%; Pred. No. 1.5e+05;
ive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Seed and Berry LLP STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                162 CGGGAGGTCCGGGAGGCGACCGTAGTCGCGCCGCCG 197
                                                                                                                                                                                                                                    2 CGGAAGGCCGGAAAGGCGAGUGAGGUCUCGCCGCAG 37
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ATTORNEY/AGENT INFORMATION:
NAME: MCMASters, David D.
REGISTRATION NUMBER: 33,963
REGISTRATION NUMBER: 920010.426C2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 324, Application US/10085906
Publication No. US20030054371A1
GENERAL INFORMATION:
APPLICANT: Ying, Vincent
APPLICANT: Wu, Paul
APPLICANT: Gray, Gary S.
                                                                                                                                                                                                                                                                                                                                                        Sequence 774, Application US/09263959
Patent No. US20020150891A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 9200
TELECOMMUNICATION INFORMATION:
TELEFONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 774:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 ACACACACACATACACACACACA 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-
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       ORGANISM: Artificial Sequence
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Matches 21; Conservative
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 6300 CCITY: Seattle
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US-10-085-906-324/c
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US-09-263-959-774
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US-09-263-959-774
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APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Draper, Ken
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of NF-Kappa B
FILE REFERENCE: 400/022 (MBHB00-812-D)
CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT FILING DATE: 2001-05-23
NUMBER OF SEQ ID NOS: 3329
SOFTWARE: Patentin version 3.0
SEQ ID NO 1826
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                                                                     'THER INFORMATION: Description of Artificial Sequence:
'OTHER INFORMATION: oligonucleotides, linkages between positions of the INFORMATION: oligonucleotides, linkages between of the INFORMATION: positions 20 to 28 are N3' to N5' OTHER INFORMATION: positions 20 to 28 are N3' to N5' of the INFORMATION: position 28 is modified by a 3'-aminodeoxyribosyl coffer INFORMATION: residue
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| Sequence 1526, Application US/09801274 |
| Patent No. US20020032319A1 |
| GENERAL INFORMATION: |
| APPLICANT: Cargill, Michele |
| APPLICANT: Ireland, James S. |
| APPLICANT: Lander, Eric S. |
| TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS FILE REFERENCE: 2525.2009-001 |
| CURRENT FILING DATE: 2001-03-07 |
| PRIOR APPLICATION NUMBER: US 60/187,510 |
| PRIOR PILING DATE: 2000-03-07 |
| PRIOR PILING DATE: 2000-03-07 |
| PRIOR FILING DATE: 2000-05-22 |
| NUMBER OF SEQ ID NOS: 1802 |
| SOFTWARE: FESTSEQ for Windows Version 4.0 |
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                                                                                                                                                                                                                                                                                    Length 28;
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                                                                                                                                                                                                                                                                                    0.4%; Score 20; DB 10; L 100.0%; Pred. No. 1.5e+05; tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1;
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                        TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                          20 GGGTGAACTGGGGGAGGATT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                           Query Match 0.4%
Best Local Similarity 100.0
Matches 20; Conservative
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LENGTH: 31
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LENGTH: 28
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Gaps

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Squence 15, Application US/09966781A

Publication No. US20030036184A1

GENERAL INFORMATION:
APPLICANT: SOULARD, PATRICIA
TITLE OF INVENTION: SELECTING COMPOUNDS WHICH INHIBIT PDE7 ENZYME ACTIVITY
FILE REPERENCE: A0000281US
CURRENT APPLICATION NUMBER: US/09/966,781A

PRIOR APPLICATION NUMBER: EP004026837

PRIOR FILING DATE: 2001-09-28

PRIOR FILING DATE: 2000-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic RNA US-09-828-034-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.4%; Score 19.4; DB 10; Length 24; Best Local Similarity 95.2%; Pred. No. 2e+05; Matches 20; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                ch 0.4%; Score 19.6; DB 9; Length 3:
1 Similarity 73.5%; Pred. No. 2.5e+05;
25; Conservative 0; Mismatches 9; Indels
                                  TITLE OF INVENTION: ALLELE SPECIFIC PCR FOR GENOTYPING
                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Synthetic oligonucleotide primer US-10-231-381-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2300 ATATAAACATCACACACACAGACAGACACACA 2333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Zhong, Weidong
APPLICANT: APONG, Weidong
APPLICANT: Hong, Zhi
APPLICANT: Hong, Zhi
TILE OF INVENTION: HCV REPLICASE COMPLEXES
CURRENT APPLICATION NUMBER: US/09/828,034
CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: U.S. 60/195,852
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PATENTIN VOI: 2.1
                                                    FILE REFERENCE: 02940201aa
CURRENT APPLICATION NUMBER: US/10/231,381
CURRENT FILING DATE: 2002-08-30
FRIOR APPLICATION NUMBER: 60/315,776
PRIOR FILING DATE: 2001-08-30
NUMBER OF SEQ ID NOS: 44
SOFTWARE: PatentIn version 3.1
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 12, Application US/09828034 Patent No. US20020064771A1
                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 CCGCCCGCCCGCCGCC 23
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SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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US-09-966-781A-15/c
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                                                                                                                                                                                                                                                                                 TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                 FEATURE:
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Sublication No. US20030096236A1
GENERAL INFORMATION:
APPLICANT: MURPHY, Patricia D.
TITLE OF INVENTION: Determining Common Functional Alleles in a Population and Uses Th
CURRENT APPLICATION NUMBER: US/09/923,327
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TITLE OF INVENTION: POLYMORPHIC ELEMENTS IN THE
TITLE OF INVENTION: COSTIMULATORY RECEPTOR LOCUS AND USES THEREOF
FILE REPERENCE: GNN-5343CP2
CURRENT APPLICATION UNBER: US/10/085,906
PRIOR APPLICATION UNBER: US/20/2027
PRIOR APPLICATION UNBER: US 09/534,061
PRIOR PILING DATE: 1099-03-25
PRIOR FILING DATE: 1000-03-24
PRIOR FILING DATE: 2000-03-24
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 545
SOFTWARE: FASSLEE OF Windows Version 4.0
LENGTHAN: 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.4%; Score 19.8; DB 9; Length 32; Best Local Similarity 77.4%; Pred. No. 1.9e+05; Matches 24; Conservative 0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.4%; Score 19.6; DB 9; Length 38; Best Local Similarity 73.5%; Pred. No. 2.4e+05; Matches 25; Conservative 0; Mismatches 9; Indels
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PRIOR APPLICATION NUMBER: US 08/598,591
PRIOR FILING DATE: 1996-02-12
PRIOR PELING DATE: 1997-02-12
PRIOR PELING DATE: 1997-02-12
PRIOR APPLICATION NUMBER: US 08/796,772
PRIOR PELING DATE: 1997-08-04
PRIOR PELING DATE: 1997-08-04
PRIOR FILING DATE: 1998-05-22
PRIOR PELING DATE: 1998-05-22
PRIOR APPLICATION NUMBER: US 09/129,134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1998-08-04
PRIOR APPLICATION NUMBER: US 09/524,794
PRIOR FILING DATE: 2000-03-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 260
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Homo sapiens
US-10-085-906-324
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; ORGANISM: Homo sapiens
US-09-923-327-60
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US-09-923-327-60
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LENGTH: 38
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194 GCCGCGCAGGACCAGGAGGAGGAG 217
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APPLICANT: Bakker, Hendrikus A.C.
APPLICANT: Bosch, Hendrik J.
                                                                                                                                                                                                                                 ; sequence 11, Application US/09772656; Patent No. US20010049832A1; GENERAL INFORMATION:
       194 GCCGCGCAGGACCAGGAGGAGGAG 217
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                                           4 GCTGCGCAGGAGCAGGAGCAGGAG 27
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SOFTWARE: Patentin version 3.1
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                                                                                                                                                                                      RESULT 42
US-09-772-656-11
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0.4%; Score 19.2; DB 10; Length 27;
Best Local Similarity 87.5%; Pred. No. 2.4e+05;
Matches 21; Conservative 0; Mismatches 3; Indels C
                                                                                                                                        ; OTHER INFORMATION: Description of Artificial Sequence:Primer US-09-966-781A-15
                                                                                                                                                                                                                                                              Length 36;
                                                                                                                                                                                                                                                                                                                                  6; Indels
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S-09-772-656-3
Sequence 3, Application US/09772656
Sequence 3, Application US/09772656
Sequence 3, Application US/09772656
Sequence 3, Application US/09772656
SERENT BERENCE: US-09-772, US
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Patent No. US20010049832A1
GENERAL INFORMATION:
APPLICANT: Bruce, Wesley B.
TITLE OF INVERTION: Root Transcriptional Factor;
FILE REFERENCE: 1167R
CURRENT APPLICATION NUMBER: US/09/772,656;
CURRENT FILING DATE: 2001-01-30
PRIOR PRICATION NUMBER: US 60/178,916
PRIOR PELING DATE: 2000-02-01
PRIOR PELING DATE: 2000-02-01
PRIOR PELING DATE: 2001-01-19
; NUMBER OF SEQ ID NOS: 12
SOSTWARE: FastERQ for Windows Version 3.0
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0.4%; Score 19.4; DB 9;
Best Local Similarity 79.3%; Pred. No. 2.6e+05;
Matches 23; Conservative 0; Mismatches 6;
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                                                                       TYPE: DNA ORGANISM: Artificial Sequence
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ORGANISM: Zea mays
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LENGTH: 27
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SEQ ID NO 15
LENGTH: 36
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1 Similarity 87.5%; Pred. No. 2.4e+05;
21; Conservative 0; Mismatches 3;
APPLICATION: Wesley B.
TITLE OF INVENTION: Root Transcriptional Factor
FILE REFERENCE: 11678
CURRENT APPLICATION NUMBER: US/09/772,656
CURRENT FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/178,916
PRIOR APPLICATION NUMBER: US 00/178,916
PRIOR FILING DATE: 2000-02-01
PRIOR FILING DATE: 2001-01-19
NUMBER OF SEQ ID NOS: 12
SOFTWARE: FastSEQ for Windows Version 3.0
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FILE REFRENCE: 05032-00019
FRIOR APPLICATION NUMBER: UST/NLOO/00774
PRIOR APPLICATION NUMBER: PCT/NLOO/00774
PRIOR APPLICATION NUMBER: EP 99203523.0
PRIOR FILING DATE: 1999-10-26
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US-10-112-653-744

| Sequence 744, Application US/10112653 |
| Publication No. US20030050268A1 |
| Sequence 744, Application US/10112653 |
| Publication No. US20030050268A1 |
| Seneral Information No. US20030050268A1 |
| APPLICANT: Krieg, Arthur M. |
| APPLICANT: Krieg, Daniel J. |
| TITLE OF INVENTION: INMUNOSTIMILATORY NUCLEIC ACID FOR TITLE OF INVENTION: TREATMENT OF NON-ALLERGIC INFLAMMATORY DISEASES |
| FILE REFERENCE: CO1039/70606(AMS) |
| CURRENT APPLICATION NUMBER: US 60/279,642 |
| PRIOR FILING DATE: 2001-03-29 |
| NUMBER OF SEQ ID NOS: 1040 |
| SOFTHARE: EastSEQ for Windows Version 3.0 |
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                                                                                                APPLICANT: Hartmann, Gunther
TITLE OF INVENTION: Methods for Enhancing Antibody-Induced
TITLE OF INVENTION: Methods for Enhancing Antibody-Induced
TITLE OF INVENTION: Cell Lyais and Treating Cancer
FILE REFERENCE: C1037/7052 (AMS)
CURRENT APPLICATION NUMBER: US/09/888,326
CURRENT APPLICATION NUMBER: US 60/213,346
PRIOR FILING DATE: 2000-06-22
PRIOR FILING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 848
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 172
LENGTH: 35
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Local Similarity 71.4%; Pred. No. 3.2e+05;
les 25; Conservative 0; Mismatches 10
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0.4%; Score 19; DB 9;
Best Local Similarity 71.4%; Pred. No. 3.2e+05;
Matches 25; Conservative 0; Mismatches 10
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Publication No. US20030055014A1
GENERAL INFORMATION:
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                                GENERAL INFORMATION:
APPLICANT: Weiner, George
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Matches
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Publication No. US200300920091
GENERAL INFORMATION:
APPLICANT: Kala Palm
TITLE OF INVENTION:
PROFILING TUNOR SPECIFIC MARKERS FOR THE
FILE REPERENCE: CEMINES.002A
CURRENT FILING DATE: 2001-11-13
PRIOR APPLICATION NUMBER: 60/249,508
PRIOR FILING DATE: 2000-11-16
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100.0%; Pred. No. 2.1e+05;
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Sequence 1, Application US/09932129

Patent No. US20020119533A1

GENERAL INFORMATION:

APPLICANT: Brown, Bob D.

TITLE OF INVENTION: AMPLIFICATION PRIMER PAIRS AND USE

TITLE OF INVENTION: THEREOF

FILE REPREBRICE: OASBIO.002C1

CURRENT PILICATION NUMBER: US/09/932,129

CURRENT FILING DATE: 2000-00-16

PRIOR APPLICATION NUMBER: PCT/US00/09230

PRIOR FILING DATE: 1999-04-08

PRIOR FILING DATE: 1999-04-08

PRIOR PRIOR APPLICATION NUMBER: US 60/128,378

PRIOR FILING DATE: 1999-04-08

SOFTWARE: FastSEQ for Windows Version 4.0

LENGTH. 10
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0.4%; Score 19; DB.
Best Local Similarity 100.0%; Pred. No. 2.1/
Matches 19; Conservative 0; Mismatches
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 377
LENGTH: 30
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Publication No. US20030026801A1
    26 TCACAAATCCTAATAGATGCATAG 3
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ORGANISM: Artificial Sequence
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US-09-888-326-172
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APPLICANT: Petersen, Yves
TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the
TITLE OF INVENTION: Treatment of Asthma and Allergy
TITLE OF INVENTION: Treatment of Asthma and Allergy
FILE REFERENCE: C1037/7013 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/09/776,479
CURRENT FILING DATE: 2001-02-02
PRIOR FILING DATE: 2001-02-03
PRIOR FILING DATE: 2000-02-03
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
SEG ID NO 771
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GENERAL INFORMATION:
APPLICANT: Brown, Bob D.
TITLE OF INVENTION: THEREOF
TITLE OF INVENTION: THEREOF
FILE REFERENCE: OASBIO.002C1
CURRENT PELLING DATE: 2001-08-16
CURRENT PELLING DATE: 2001-08-16
PRIOR APPLICATION NUMBER: PCT/USO0/09230
PRIOR FILING DATE: 2000-04-07
PRIOR FILING DATE: 1999-04-08
PRIOR FILING DATE: 1999-04-08
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                            Query Match 0.4%; Score 19; DB 9; Length 35; Best Local Similarity 71.4%; Pred. No. 3.2e+05; Matches 25; Conservative 0; Mismatches 10; Indels
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CURRENT FILING DATE: 2001-12-18
PRIOR APPLICATION NUMBER: US 60/255,534
PRIOR FILING DATE: 2000-12-14
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 771
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                                                                                                                                                                                                                                                                 OTHER INFORMATION: Synthetic Sequence US-10-017-995-771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 771, Application US/09776479; Sequence 771, Application US/097087848A1; GENERAL INFORMATION: APPLICANT: Bratzler, Robert L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 4, Application US/09932129
; Patent No. US20020119533A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                           TYPE: DNA ORGANISM: Artificial Sequence
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US-09-932-129-4
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SECTWARE: FastSEQ for Windows Version 4.0

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/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
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GenCore version 5.1.5
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-07-936-421-15
US-08-217-082A-2
US-07-936-421-20
US-09-080-285-2
US-09-080-285-2
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US-09-233-341-8 US-09-253-341-8 US-09-337-635-20 US-09-261-040-8 US-09-337-280-20 US-09-337-280-20 US-08-311-342-12 US-08-117-981-11	US-09-306-290-2 US-09-453-195A-10 US-09-538-700-1236 PCT-US93-02142-11 US-08-217-082A-17	US-08-894-784-14 US-08-894-784-14 US-08-894-784-35 US-08-894-784-35 US-09-080-285-17 US-09-080-285-24	US-09-110-22-5 US-09-030-701-27 US-09-286-098-59 US-09-286-098-104 US-09-078-954-14	US-09-724-426-1/ US-09-724-426-24 US-09-191-170-53 US-08-063-167A-59 US-08-440-740A-59 US-08-344-155C-59 US-08-344-155C-59	US-v8-y1-2-2-3-3-3-3-3-3-3-3-3-3-3-3-3-3-3-3-3-	US-08-927-561-42 PCT-US96-09388-42 US-08-504-511A-6 US-08-467-126-6 US-08-629-001A-44 US-08-642-712-5 US-09-411-291-5 US-09-268-505B-13 US-09-268-505B-13	US-08-433-330-118 US-08-435-350-3 US-07-915-245-1 US-08-145-704-2 US-08-145-704-4 US-08-987-574-2 US-08-535-168-2 US-08-535-168-2 US-08-535-168-4 US-09-017-974-4
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Sequence 28, Appl Sequence 1008, Appl Sequence 25, Appl Sequence 25, Appl Sequence 25, Appl Sequence 31, Appl Sequence 60, Appl Sequence 10, Appl Sequence 10, Appl Sequence 11, Appl Sequence 1	e uoue
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Sequence 4, Appli Sequence 2, Appli Sequence 10, Appli Sequence 11, Appli Sequence 11, Appli Sequence 11, Appli Sequence 11, Appli Sequence 12, Appli Sequence 14, Appli Sequence 14, Appli Sequence 16, Appli Sequence 17, Appli Sequence 16, Appli Sequence 17, Appli Sequence 16, Appli Sequence 17, Appli Sequence 17, Appli Sequence 16, Appli Sequence 17, Appli Sequence 17, Appli Sequence 11, Appli Sequence 12, Appli Sequence 12, Appli Sequence 12, Appli Sequence 11, Appli Sequence 12, Appli Sequence 11, Appli Sequence 12, Appli Sequence 12, Appli Sequence 12, Appli Sequence 13, Appli Sequence 14, Appli Sequence 17, Appli	
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GENERAL INFORMATION:
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US-08-217-082A-2
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US-08-217-082A-2
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                                                                                                                                                                                                                                                                                                                          Query Match
0.7%; Score 37; DB 1; Length 37;
Best Local Similarity 70.3%; Pred. No. 2.6;
Matches 26; Conservative 11; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. DOS (Version 5.0) SOFTWARE: WORGPERFECT (Version 5.1) CURRENT APPLICATION DATA: APPLICATION NUMBER: US/07/936,421 FILING DATE: 19920826 CLASSIFICATION: 435 PRIOR APPLICATION DATA: including application PRIOR APPLICATION DATA: described below: APPLICATION DATA: described below: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: James D. Thompson
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: TREATMENT OF DISEASES CAUSED
TITLE OF INVENTION: BY EXPRESSION OF THE BCL-2
TITLE OF INVENTION: GENE
                                                                                                                                                                                                                                                                                                                                                                                                          1415 GGCGAGAGGTGCCGTTGCCCCGTTGCTTTCCTCT 1451
                                                                                                                                                                                                                                                                                                                                                                                                                               1 GGCGAGAGGGCCGUUGGCCCCGUUGCUUUCCUCU 37
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TELECOMMUNICATION INFORMATION:
TELEPAX: (213) 955-0440
IELEX: 67-3510
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 15, Application US/07936421
Patent No. 5750390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Lyon & Lyon
611 West Sixth Street
                   NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
                                                                                            TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      32,327
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                      TYPE: NUCLEIC ACID
STRANDEDNESS: single
TOPOLOGY: linear
US-07-936-421-2
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MEDIUM TYPE: 3.5" DI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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California
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US-07-936-421-15
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US-07-936-421-15
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APPLICANT: Reed, John
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES FOR INHIBITING THE
TITLE OF INVENTION: ARMYH OF CELLS EXPRESSING THE HUMAN BCL-2 GENE
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
ADDRESSEE: P.C.
                                                 Gaps
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  Length 36;
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                                              Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPPRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/217,082A
Score 36; DB 1;
Pred. No. 4.5;
5; Mismatches
                                                                                    1997 ACCIGAACCGCACCTGCACCTGGATCCAGGATA 2032
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100.0%; Pred. No. 7.8;
tive 0; Mismatches
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APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/288,692
FILING DATE: 22-DEC-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 24-MAR-1994
CLASSIFICATION: 435
(IOR ADDITOR: 435
                                                                                                                                                                                                                                    ; Sequence 2, Application US/08217082A; Patent No. 5734033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 4
US-07-936-421-20
; Sequence 20, Application US/07936421
; Patent No. 5750390
                                                                                                                                                                                                                                                                                                                                                                                                                                                    224 Airport Parkway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Fortney, Andrew D.
REGISTRAITON NUMBER: 34,600
REFERENCE/DOCKET NUMBER: 33
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (408) 436-2070
TELEPAX: (408) 436-2075
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 35 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA (genomic)
Query Match 0.7%;
Best Local Similarity 86.1%;
Matches 31; Conservative
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Matches 35; Conservative
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STRANDEDNESS: single
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CITY: San Jose
STATE: California
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Sequence 2. 4 Application US/09080285

Patent No. 6040181

GENERAL INFORMATION:

APPLICANT: Reed, John
TITLE OF INVENTION:

NUMBER OF SEQUENCES: 29

CORRESPONDENCE ADDRESS:

ADDRESSEE: P.C.
STREET: 1755 S. Jefferson Davis Hwy., Suite 400
CITY: Arlington
STATE: Virginia

COUNTY: N. O.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.7%; Score 35; DB 2; Length 35; Best Local Similarity 100.0%; Pred. No. 7.8; Matches 35; Conservative 0; Mismatches 0; Indels
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PatentIn Release #1.0, Version #1.25
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                                                                                                                PRIOR APPLICATION DATA

APPLICATION NUMBER: US 08/124,256
FILING DATE: 20-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/840,716
FILING DATE: 21-FEB-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/288,692
FILING DATE: 22-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: FORTHREY, ANGINEW:
NAME: FORTHREY, ANGINEW:
NAME: FORTHREY, ANGINEW:
NAME: FORTHREY, ANGINER: 34,600
REGISTRATION NUMBER: 34,600
REGISTRATION NUMBER: 34,600
RECISTRATION NUMBER: 335-070-5
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 35 base pairs
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APPLICATION NUMBER: US 08/465,485
FILING DATE: 05-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/124,256
FILING DATE: 20-58E-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/840,716
                                 OMBER: US/08/465,485A
05-JUN-1995
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA (genomic)
                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 05-JUN-199
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
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MOLECULE TYPE: I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ANTI-SENSE:
US-08-465-485A-2
       SOFTWARE:
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Patent No. 581066
BAREAL INFORMATION:
APPLICANT: Reed, John
TITLE OF INVENTION: Regulation of bcl-2 Gene Expression
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
ADDRESSEE: P.C.
STREET: 1755 S. Jefferson Davis Hwy., Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match

0.7%; Score 35; DB 1; Length 35;
Best Local Similarity 85.7%; Pred. No. 7.8;
Matches 30; Conservative 5; Mismatches 0; Indels
                                                     TITLE OF INVENTION: METHOD AND REAGENT FOR TITLE OF INVENTION: TREATMENT OF DISEASES CAUSED TITLE OF INVENTION: BY EXPRESSION OF THE BCL-2 TITLE OF INVENTION: GENE
                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NAMER: US/07/936,421
FILING DATE: 19920826
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM Compatible
COMPUTER: IBM COmpatible
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PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
FILING DATE:
FILING DATE:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 611 West Sixth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
                                 Kenneth G. Draper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS: single
                                                                                                                                                      NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                       CITY: Los Angeles
STATE: California
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STATE: Virginia
COUNTRY: U.S.A.
ZIP: 22202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
US-07-936-421-20
                                                                                                                                                                                                                                                                                                       USA
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US-08-465-485A-2
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskett
COMPUTER: IBM COmpatible
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Best Local Similarity 76.5%
Matches 26; Conservative
                                                                                                                                                               CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
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CITY: Los Angeles
STATE: California
COUNTRY: USA
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OPERATING SYSTEM:
    California
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US-07-936-421-21
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US-09-724-426-2
Sequence 2, Application US/09724426
Sequence 2, Application US/09724426
Patent No. 6414134
GENERAL INFORMATION:
TITLE OF INVENTION: Regulation of BCL-2 Gene Expression
TITLE OF INVENTION: Regulation of BCL-2 Gene Expression
CURRENT FILING DATE: 2000-11-28
UNRENT FILING DATE: 2000-11-28
SOFTRANE: PatentIn version 3.0
SEQ. ID NO. 2
                                                                                                                                                                                                                                                                                                                                                                                                                             0.7%; Score 35; DB 3; Length 35; 100.0%; Pred. No. 7.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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APPLICANT: Annew D. Thompson
AITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: TREATMENT OF DISEASES CAUSED
TITLE OF INVENTION: BY EXPRESSION OF THE BCL-2
TITLE OF INVENTION:
GENE
NUMBER OF SEQUENCES: 22
FILING DATE: 21-FEB-1992
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/288,692
FILING DATE: 22-DEC-1988
ATTORNEY/AGENT INFORMATION:
NAME: FOATHREY, AND THE W. D.
REGISTRATION NUMBER: 34,600
REFERENCE/DOCKET NUMBER: 3335-070-55 CONT TELECHONICATION INFORMATION:
TELEPHONE: (408) 436-2075
INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS:
LENGTH: 35 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1442 CTTTTCCTCTGGGAAGGATGGCGCACGCTGGGAGA 1476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1442 CTTTTCCTCTGGGAAGGATGGCGCACGCTGGGAGA 1476
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3: Lyon & Lyon
611 West Sixth Street
                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           35; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        35; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Los Angeles
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US-07-936-421-16
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Gaps
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APPLICANT: James D. Thompson
APPLICANT: James D. Thompson
APPLICANT: TRENENT G. Draper
TITLE OF INVENTION: TREATMENT OF DISEASES CAUSED
TITLE OF INVENTION: BY EXPRESSION OF THE BCL-2
TITLE OF INVENTION: GENE
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESSE:
ADDRESSEE: Lyon & Lyon
STREET: 611 West Sixth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SYSTEM: IBM P.C. DOS (Version 5.0) WordPerfect (Version 5.1)
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
                                                                                                                                                                     FILING DATE: 19920826
CLASSIFTCATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
APPLICATION NUMBER:
APPLICATION NUMBER:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3.5" Diskette, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            including application described below:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2043 GGATGCCTTTGTGGAACTGTACGGCCCCAGCATG 2076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GGAUGCCUUUGUGGAACUGUACGCCCCAGCAUG 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/936,421
FILING DATE: 19920826
CLASSIFICATION: 435
                                                                                                                                                APPLICATION NUMBER: US/07/936,421 FILING DATE: 19920826
                                                                                                                                                                                                                                                                                                                                                                                                                        197/243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21, Application US/07936421
Patent No. 5750390
                                                                                                                                                                                                                                                                                                                                                                       NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: (7-3510
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
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Gaps

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APPLICANT: Capaccioli, Sergio
APPLICANT: Morelli, Susanna
APPLICANT: Nicolin, Angelo
TITLE OF INVENTION: CELLS HAVING A T(14:18) TRANSLOCATION AND
TITLE OF INVENTION: CELLS HAVING A T(14:18) TRANSLOCATION AND
TITLE OF INVENTION: CELLS HAVING A T(14:18) TRANSLOCATION AND
TITLE OF INVENTION: TRAIMENT OF SAID TUMOR CELLS
NUMBER OF SEQUENCES: 49
ADDRESSEE: FINNEGAN, HENDERSON, FARABOW, GARRETT &
ADDRESSEE: DUNNER, LLP
                                                                                                                          ó
                                                                          Score 33; DB 1; Length 33;
Pred. No. 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.6%; Score 33; DB 3; Length 33;
100.0%; Pred. No. 24;
Live 0; Mismatches 0; Indels
                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COmpatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05999.0005-00000
                                                                                                                                                                   1592 CACCGGGCATCTTCTCTCCCCAGCCCGGGCACA 1624
                                                                                                                     5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4289 GCAATTCCGCATTTAATTCATGGTATTCAGGAT 4321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/894,784
FILING DATE: 15-DEC-1997
FILING DATE: 15-DEC-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP96/00852
FILING DATE: 02-MAR-1996
PRIOR APPLICATION NUMBER: IL M195 A 000420
FILING DATE: 03-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Paul, John C.
REGISTRATION NUMBER: 30,413
                                                                                                                                                                                           1 CACCGGGCAUCUUCUCCUCCCAGCCCGGGCACA 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GCAATTCCGCATTTAATTCATGGTATTCAGGAT 33
                                                                                                                                                                                                                                                                                                                                Sequence 44, Application US/08894784 Patent No. 6005095 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 05
TELECOMUNICATION INFORMATION:
TELEPHONE: 202-408-4000
TELEFAX: 202-408-4400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
                                                                        Query Match 0.6%;
Best Local Similarity 84.8%;
Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 1300 I Street, NW CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; MOLECULE TYPE: DNA
US-08-894-784-44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 20005-3315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 33; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                        US-08-894-784-44
              ;
US-07-936-421-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE:
                                                                                                                                                                                                                                                                                       RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                 Ouery Match 0.7%; Score 34; DB 1; Length 34; Best Local Similarity 73.5%; Pred. No. 14; Matches 25; Conservative 9; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: James D. Thompson
APPLICANT: James D. Thompson
APPLICANT: Renneth G. Draper
TITLE OF INVENTION: TREATMENT OF DISEASES CAUSED
TITLE OF INVENTION: BY EXPRESSION OF THE BCL-2
TITLE OF INVENTION: GENE
TITLE OF INVENTION: GENE
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 611 West Sixth Street
CITY: Los Angeles
STATE: Galifornia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. DOS (Version 5.0) SOFTWARE: WOLDERFECT (Version 5.1) CURRENT APPLICATION DATA: APPLICATION NUMBER: US/07/936,421 FILING DATE: 19920826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2235 AATATGCATTGTCAGTGATGTACCATGAAACAAA 2268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 AAUAUGCAUUGUCAGUGAACCAUGAAACAAA 34
                                    ATTORNEY AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 197/243
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5, Application US/07936421
Patent No. 5750390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 19'
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                           LENGTH: 34
TYPE: NUCLEIC ACID
STRANDEDNESS: Single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            single
APPLICATION NUMBER:
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                   FILING DATE:
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US-07-936-421-5
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Gaps

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Sequence 415, Application US/08222177A
Patent No. 5582979
GENERAL INFORMATION:
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Capaccioli, Sergio
APPLICANT: Morelli, Susanna
APPLICANT: Morelli, Susanna
APPLICANT: Morelli, Susanna
APPLICANT: Micolin, Angelo
TITLE OF INVENTION: CELLS HAVING A T(14:18) TRANSCOIATED TO TUMOR
TITLE OF INVENTION: OLIGODEOXYNUCLEOTIDES USEFUL IN THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF SAID TUMOR CELLS
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: FINNEGAM, HENDERSON, FARABOW, GARRETT &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.6%; Score 30; DB 1; L. Best Local Similarity 100.0%; Pred. No. 1.3e+02; Matches 30; Conservative 0; Mismatches 0;
                                                                                                                                                                                                           ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2310 CACACACACAGACAGACACACACACACA 2339
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 07/341,562 FILING DATE: 21-APR-1989 ATTORNEY/AGENT INFORMATION: NAME: Sara, Charles S. REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-894-784-19; Sequence 19, Application US/08894784; Patent No. 6005095; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 415:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DNA (genomic)
US-08-222-177A-415
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Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                             ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                      Wisconsin : USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                                                                                                                                                                                                                               Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
US-08-222-177A-415
                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 13
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GENERAL INFORMATION:
APPLICANT: Capaccioli, Sergio
APPLICANT: Mocalli, Susanna
APPLICANT: Miccalin, Annelio
TITLE OF INVENTION: ANTISENSE TRANSCRIPT ASSOCIATED TO TUMOR
TITLE OF INVENTION: CELLS HAVING A T(14:18) TRANSLOCATION AND
TITLE OF INVENTION: OLIGODEOXYNOCLEOTIDES USEFUL IN THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF SAID TUMOR CELLS
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                   SOFTWARES PATENTING SISLEM:
CURRENT APPLICATION DATA:
APPLICATION NUMBER:
TLING DATE: 15-DEC-1997
FILING DATE: 15-DEC-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT_EP96/00852
FILING DATE: 02-MAR-1996
FILING DATE: 02-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1L MI95 A 000420
FILING DATE: 03-MAR-1996
ATTORNEY/AGENT INFORMATION:
NAME: PAUL, JOHN C.
REGISTRATION NUMBER: 05999.0005-0000
TELEPROCMONICATION INFORMATION:
TELEPROMONICATION INFORMATION:
TELEPRAX: 202-408-4400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: FINNEGAN, HENDERSON, FARABOW, GARRETT ADDRESSEE: DUNNER, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,008/094,784
FILING DATE: 15-DEC-1997
PRIOR APPLICATION DATA:
FILING DATE: 02-MAR-1996
PRIOR DATE: 02-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 3; Le . 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.6%; Score 30; DB Best Local Similarity 100.0%; Pred. No. 1.3 Matches 30; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4256 GACCTIGITICITGAAGGITICCTCGTCCC 4285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GACCTTGTTTCTTGAAGGTTTCCTCGTCCC 30
                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-894-784-23
; Sequence 23, Application US/08894784
; Patent No. 6005095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 1300 I Street, NW CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: linear; MOLECULE TYPE: DNA US-08-894-784-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 20005-3315
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APPLICANT: Capacacioli, Sergio
APPLICANT: Morelli, Susanna
TITLE OF INVENTION: CELLS HAVING A T(14:18) TRANSCORTION AND
TITLE OF INVENTION: OLIGODEOXYNICLEOTIDES USEFUL IN THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF SAID TUMOR CELLS
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                    Length 28;
                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: RIOPPY disk
MEDIUM TYPE: RIOPPY disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: DATE: RIOPPY disk
COMPUTER: DATE: PC COMPATIBLE
COMPUTER: STATEM: PC COMPATIBLE
COMPUTER: DATE: DC COMPATIBLE
SOUTHARE: PRECHIN Release #1.0, Version #1.30
CURRENT APPLICATION DATE: US/08/894,784
FILING DATE: 15-DEC-1997
PRIOR APPLICATION NUMBER: PCT/EP96/00852
FILING DATE: 02-MAR.1996
PILING APPLICATION NUMBER: IL MI95 A 000420
FILING DATE: 03-MAR.1995
ATTORNEY/AGENT INFORMATION:
NAME: PRUL, JOHN C.
REGISTRATION NUMBER: 30,413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: FINNEGAN, HENDERSON, FARABOW, GARRETT ADDRESSEE: DUNNER, LLP
                                                                  Score 28; DB 1; Pred. No. 3.9e+02; 5; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.6%; Score 28; DB 3; Best Local Similarity 100.0%; Pred. No. 4e+02; Matches 28; Conservative 0; Mismatches C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30,413
BER: 05999.0005-00000
                                                                                                                                                           1729 CCTGTGGTCCACCTGGCCCTCCGCCAAG 1756
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                                                                                                                                                                                                                                                                                                                Sequence 20, Application US/08894784 Patent No. 6005095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMUNICATION INFORMATION:
TELECHMUNICATION INFORMATION:
TELEPHONE: 202-408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JUNNER, LLP UNNER, LLP UNINEET: 1300 I Street, NW CITY: Washington
                                                                  Query Match 0.6%;
Best Local Similarity 82.1%;
Matches 23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
  TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                          RESULT 16
US-08-894-784-20/c
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           ;
US-07-936-421-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 17
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OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/936,421
FILING DATE: 19920826
CLASSIFICATION A35
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                      1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: James D. Thompson
APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: TREATMENT OF DISEASES CAUSI
TITLE OF INVENTION: BY EXPRESSION OF THE BCL-2
TITLE OF INVENTION: GENE
NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.6%; Score 30; DB 3; Best Local Similarity 100.0%; Pred. No. 1.3e+0 Matches 30; Conservative 0; Mismatches
                     FILING DATE: 03-WAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Paul, John C.
REGISTRATION NUMBER: 30,413
REFERENCE/DOCKET NUMBER: 05999.0005-00000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4256 GACCTIGITICITGAAGGTITCCTCGTCCC 4285
IL M195 A 000420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 611 West Sixth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/07936421 Patent No. 5750390 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327 REFERENCE/DOCKET NUMBER: 19 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                 TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELERX: 67-3510
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
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STRANDEDNESS: single
    APPLICATION NUMBER:
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CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: DNA
-08-894-784-23
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us-09-375-514-19.rni

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FILING DATE:
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Patent No. 5582979
GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS: ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                        APPLICANT: Capaccioli, Sergio
APPLICANT: Morelli, Susanna
APPLICANT: Mocelli, Susanna
APPLICANT: Mocelli, Susanna
APPLICANT: Mocelli, Susanna
TITLE OF INVENTION: CELLS HAVING A T(14:18) TRANSLOCATION AND
TITLE OF INVENTION: CLIGODECOXYNCLEOTIDES USEFUL IN THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF SAID TUMOR CELLS
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.6%; Score 28; DB 3; Length 30;
100.0%; Pred. No. 4e+02;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTEK: LEM PC COMPGALLAGE
CORENTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PACEDIIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/894,784
FILING DATE: 15-DEC-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/FP96/00852
FILING APPLICATION DATA:
APPLICATION NUMBER: IL MI95 A 000420
FILING APPLICATION NUMBER: IL MI95 A 000420
FILING APPLICATION NUMBER: IL MI95 A 000420
FILING APPLICATION NUMBER: 30.413
REFERRATION NUMBER: 30,413
REFERRATION NUMBER: 30,413
                                                                                                                                                                                                                                                                                                      ADDRESSEE: FINNEGAN, HENDERSON, FARABOW, GARRETT ADDRESSEE: DUNNER, LLP STREET: 1300 I Street, NW CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4504 CTTCAGGGTCTTCCTGAAATGCAGTGGT 4531
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US-08-894-784-24/c
; Sequence 24, Application US/08894784
; Patent No. 6005095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 24: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 30 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Madison
STATE: Wisconsin
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Matches 28; Conserv
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                                                                       GENERAL INFORMATION:
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Sequence 403, Application US/08222177A
Patent No. 5882979
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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80.0%; Pred. No. 7.6e+02;
tive 0; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2310 CACACACACAGACAGACACACACACACAACAATTAACA 2349
                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
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SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPELCATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATE: 05/341,562
FILING DATE: 21-APR-1989
ATORNEY/AGENT INFORMATION:
NAME: Sara, Challes S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                   09865.601
                                                                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: SAIA, CHAILES S.
REGISTRATION NUMBER: 30,492
                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 8000 Excelsior Drive, CITY: Madison STATE: Wisconsin COUNTR: USA ZIP: 53717-1914 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPATEN: OPERATING SYSTEM: PC-DOS/MS-DC
                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 09:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 175:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfd42rs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.5%
Best Local Similarity 80.0%
Matches 32; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 40 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
53717-1914
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Gaps

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Pred. No. 6.8e+02; 3; Mismatches 0;

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2081 CICIGITIGATITCICCIGGCIGICIC 2107
                                                                                     ilarity 51.9%; Pr
Conservative 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 31 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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Matches 28; Conservative
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  Best Local Similarity
Matches 14; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Madison
STATE: Wisconsin
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; CLONE: mfd57rs
US-08-222-177A-215
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US-08-222-177A-235
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                                                                                                                                                                                                                                                                                      Length 40;
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APPLICANT: Kenneth G. Draper
TITLE OF INVENTION: METHOD AND REAGENT FOR
TITLE OF INVENTION: TREATMENT OF DISEASES CAUSED
TITLE OF INVENTION: GENE
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 611 West Sixth Street
                                                                                                                                                                                                                                                                                                                                                                 2310 CACACACACAGACAGACACACACACACAAATTAACA 2349
                                                                                                                                                                                                                                                                                                                                                                                      1 CACACACACACACACACACACACACACACACACACA 40
                                                                                                                                                                                                                                                                                  Query Match 0.5%; Score 27.2; DB 1; Best Local Similarity 80.0%; Pred. No. 7.6e+02; Matches 32; Conservative 0; Mismatches 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: 1BM Compatible
OPERATING SYSTEM: 1BM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below:
  09865.601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/07/936,421
FILING DATE: 19920826
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Patent No. 5750390
                                                                                                    403:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327 REFERENCE/DOCKET NUMBER: 19 TELECOMMUNICATION INFORMATION:
REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION
TELEPHONE: (608) 831-2106
TELEFAX: (608) 831-2106
                                                                                                                                                                               STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
US-08-222-177A-403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELER: 67-3510
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 40 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-07-936-421-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-07-936-421-17
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0.5%; Score 27; DB 1; Length 27;

Query Match

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RESULT 21
US-08-222-177A-215
US-08-222-177A-215
Sequence 215, Application US/08222177A
Patent No. 5582979
GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS. OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dewlitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dc-da)n.(dg-dt)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.5%; Score 26.8; DB 1; Length 31; 93.3%; Pred. No. 8.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/222,177A FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
RILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: SATA, ChATLES S.
REFERENCE/DOCKET NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 CACACACACACACACACACACACACACA 30
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                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
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LENGTH:
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Patent No. 5582979
GENERAL INFORMATION:
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Lab PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                               ZIP: 53717-1914

ZIP: COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN RELEASE #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 26.8; DB 1;
Pred. No. 8.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
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              E: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
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                                                                                                                                                                                                                                                                                                PLICASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 CACACACACACACACACACACACACACACA 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 235: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA (genomic); IMMEDIATE SOURCE; CLONE: mfd64rs
US-08-222-177A-235
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Best Local Similarity 93.3%;
Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 31 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
CORRESPONDENCE ADDRESS:
                                                                                 Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Madison
Wisconsin
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                                                         Madison
                                                                                                    USA
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                                                                                                                                                                                                                                                                                     FILING DATE
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                                                                                                COUNTRY:
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GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dG-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
ADDRESSE: DeWitt Ross & Stevens, S.C.
STREPT: 8000 Excelsior Drive, Suite 401
CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.5%; Score 26.8; DB 1; Length 31; Best Local Similarity 93.3%; Pred. No. 8.2e+02; Matches 28; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2310 CACACACACAGACAGACACACACACACACA 2339
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
US 07/341,562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 24
2-08-222-177A-210
Sequence 210, Application US/0822177A
Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                               NAME: Sara, Charles S.
REGISTRATION NUMBER: 30, 492
REFERENCE/FOCKET NUMBER: 0986
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Sara, Charles S. REGISTRATION NUMBER: 30,492
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TELECOMMONICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                   271:
                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA (genomic); IMMEDIATE SOURCE:
CLONE: mfd79rs
US-08-222-117A-271
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MOLECULE TYPE: DNA (genomic)
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TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                 FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 31 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: double
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                                                                                                                                                                                                                                                                                                                                                      linear
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Query Match
Best Local Similarity
                                                                                                                                                   CITY: Madison
STATE: Wisconsin
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STATE: Wisconsin
   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                      FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 27
US-08-222-177A-218
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TELEX:
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APPLICANT: Weber, James L.

TITLE OF INVENTION: (dc-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
TITLE OF INVENTION: (dc-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460

CORRESPONDENCE ADDRESS:
ADDRESSEE: DGMILT ROSS & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
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                                                                        Length 32;
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                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                  Score 26.8; DB 1;
Pred. No. 8.4e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AFLILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTONREY/ABGHT INFORMATION:
NAME: SATA, Charles S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                 2310 CACACACAGAGAGACACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2310 CACACACACAGACAGACACACACACACACA 2339
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                                                                                                                                                                                                                                                                         Sequence 376, Application US/08222177A Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 26
US-08-222-177A-59
'Sequence 59, Application US/08222177A
'Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 376:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                        0.58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfdll9rs
US-08-222-177A-376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Floppy disk
                                                                    Query Match
Best Local Similarity 93.33
Matches 28, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Wisconsin
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
; IMMEDIATE SOURCE:
; CLONE: mfd55rs
US-08-222-177A-210
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Madison
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US-08-222-177A-376
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APPLICANT: Weber, James L.

TITLE OF INVENTION: LENGTH POLYMORPHISMS IN

TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Weber, James L.
TILLE OF INVENTION: LENGTH POLYMORPHISMS IN
TILLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dewlit Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/222,177A
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COMPUTER: IBM PC Compatible
OPERATUG SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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93.3%; Pred. No. 8.5e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                 ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY.AGENT INFORMATION:
NAME: Sara. Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09665.601
TELECHOME: (608) 831-2100
TELEFAX: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                            STAIL:
COUNTRY: USA
ZIP: 5317-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
--writter: IBM PC compatible
--writter: Floppy STAIL FORM:
--writter: Floppy Disk
--writter: Floppy
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IMMEDIATE SOURCE:
CLONE: mfd3rs
US-08-222-1774-59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acid
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ZIP: 53717-1914
COMPUTER READABLE FORM:
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34 base pairs

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LENGIH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Patent No. 5582979
FARENT NO. 5582979
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dG-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASCIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: SATA, Charles S.
REGISTRATION NUMBER: 30,492
REFERNICE/DOCKET NUMBER: 09865.601
TELEPHONE: (608) 831-2100
                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 218:
SEQUENCE CHARACTERISTICS:
LENGTH: 33 base pairs
TENGTH: 33 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-
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ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; CLONE: mfd58rs
US-08-222-177A-218
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                                              FILING DATE:
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APPLICATE: WEBER, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
ADDRESSE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
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                                                                                                                                                             0.5%; Score 26.8; DB 1; Length 34; 93.3%; Pred. No. 8.7e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.5%; Score 26.8; DB 1; Length 34; 93.3%; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
                                                                                                                                                                                                                                                                       2 CACACACACACACACACACACACACACA 31
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                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 172, Application US/08222177A
; Patent No. 5582979
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                     MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfd20rs
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                                                                                                                                                                                  Best Local Similarity 93.3
Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 34 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TYPE: nucleic acid
STRANDEDNESS: double
                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wisconsin
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nes 28; Conserv
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CLONE: mfd41rs
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MOLECULE TYPE:
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US-08-222-177A-110
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                                        TOPOLOGY:
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                                                                                                                                                               Query Match
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us-09-375-514-19.rni

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COMPUTER READABLE FORM:
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                                      98-222-177a-180

Fatent No. 558297108

Sequence 180, Application US/08222177A

Patent No. 55829710N:
GENERAL INFORMATION:
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460

CORRESPONDENCE ADDRESS:
ADDRESSEE: Dewlit Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 192, Application US/08222177A
Patent No. 5582979
GENERAL INFORMATION:
APPLICANT: Weber, James L.
APPLICANT: Weber, James L.
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dewitt Ross & Stevens, S.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DAYE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DAYE: 21-APR-1989
ATTORNEY-AGENT INFORMATION:
NAME: SATA, Charles S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2310 CACACACAGACAGACACACACACACACA 2339
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                                                                                                                                                                                                                                                                                              CITY: Madison
STATE: Wisconsin
COUNTRY: USA
2IP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
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INFORMATION FOR SEQ ID NO: 180:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfd44rs
US-08-222-177A-180
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TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 34 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wisconsin
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US-08-222-177A-322
Sequence 322, Application US/08222177A
Sequence 3222177A
Sequence 322217A
Sequence 3222217A
Sequence 32
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
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                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/POCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: SAIA. CHAILSS S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 CACACACACACACACACACACACACA 31
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ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPALIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 192:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfd48r8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 34 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Madison
STATE: Wisconsin
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2310 CACACACACAGACAGACACACACACACA 2339
                                                 1 CACACACACACACACACACACACACACACACA 30
                                                                                                                                                                                                                                                                                                                                                                                      Wisconsin
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                                                                                                                            RESULT 34
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TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
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                                                                                                                                                                                                                                                                             Query Match 0.5%; Score 26.8; DB 1; Length 34; Best Local Similarity 93.3%; Pred. No. 8.7e+02; Matches 28; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ch 0.5%; Score 26.8; DB 1; Length 35; 1 Similarity 93.3%; Pred. No. 8.8e+02; 28; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                            2310 CACACACAGACAGACACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                              2 CACACACACACACACACACACACACA 31
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTOKNEY/AGENT INFORMATION:
NAME: SAIA, CHAILES S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 77, Application US/08222177A Patent No. 5582979 GENERAL INFORMATION:
                                                              322:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30,492
                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfd101rs
US-08-222-177A-322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 09
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; CLONE: mfd9rs
US-08-222-1177A-77
    TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                       INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 34 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 35 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wisconsin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Madison
STATE: Wiscons
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TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-222-177A-77
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Gaps

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APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dG-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-222-177A-104

Sequence 104, Application US/08222177A

Sequence 104, Application US/08222177A

GENERAL INCEMPTION:
GENERAL INFORMATION:
APPLICANT: Weber, James L.

TITLE OF INVENTION: (dG-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME TITLE OF SEQUENCES: 460

CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                                                                                                                                                                                                                                                                                                                                  ZIP: 53717-1914

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 26.8; DB 1;
Pred. No. 8.8e+02;
0; Mismatches 2;
                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2310 CACACACAGACAGACACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASTICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: SATA, Charles S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 CACACACACACACACACACACACACACACA 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09865.601
US-08-222-177A-189
Sequence 189, Application US/08222177A
Patent No. 5582979
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA (genomic) IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.5%;
Best Local Similarity 93.3%;
Matches 28; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  35 base pairs
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RESULT 37
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APPLICANT: Weber, James L.
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
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9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.5%; Scur.
93.3%; Pred. No. >-
... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: DeWitt Ross & Stevens, S.C. 8000 Excelsior Drive, Suite 401
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: SATA, ChATLES S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 CACACACACACACACACACACACACACACA 31
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STATE: Wisconsin
COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,1778
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
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US-08-222-177A-358
Sequence 358, Application US/08222177A
, Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 104:
SEQUENCE CHARACTERISTICS:
LENGTH: 36 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfdl8rs
US-08-222-177A-104
                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                  Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 93.39
                                                        COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
                                     Wisconsin
                  Madison
                                                                                                                  MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                      FILING DATE:
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US-08-222-177A-140

Sequence 140, Application US/08222177A

Patent No. 5582979

Patent No. 5582979

Patent No. 5582979

TITLE OF INVENTION: LENGTH POLYMORPHISMS IN

TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.5%; Score 26.8; DB 1; Length 36; Best Local Similarity 93.3%; Pred. No. 9e+02; Matches 28; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2310 CACACACACAGACAGACACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 CACACACACACACACACACACACA 31
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ER: 09865.601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNBER: 20 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles 30,492
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION UNDER: 30,492
REFERENCE/DOCKET NUMBER: 0986:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2106
TELERAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfdll3rs
US-08-222-177A-358
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MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfd30Ts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 37 base pairs
TYPE: nucleic acid
STRANDENNESS: double
                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 36 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 53717-1914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDAL.
STREET: BULL
TTY: Madison
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us-09-375-514-19.rni

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Sequence 397, Application US/08222177A
Patent No. 582879
BAPPLICANT NO. 582879
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (4C-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDED ADDRESS:
ADDRESSEE: Dewitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
LENGTH POLYMORPHISMS IN (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                     COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURREMY APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.5%; Score 26.8; DB 1;
93.3%; Pred. No. 9.3e+02;
tive 0; Mismatches 2;
                                                                                                     3: DeWitt Ross & Stevens, S.C. 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2310 CACACACACAGACAGACACACACACACA 2339
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Sare, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: DNA (genomic) IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 53117-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 93.39
Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 38 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid
EDNESS: double
                     TITLE OF INVENTION: LEI
TITLE OF INVENTION: (dc
NUMBER OF SEQUENCES: 4(
CORRESPONDENCE ADDRESS:
                                                                                                                                                CITY: Madison
STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Madison
STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; CLONE: mfd50rs
US-08-222-177A-198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-222-177A-397
                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
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Sequence 68, Application US/08222177A

Sequence 68, Application US/08222177A

GENERAL IN. 5582979

GENERAL INFORMATION:

APPLICANT: Weber, James L.

TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460

CORRESPONDENCE ADDRESS:

ADDRESSEE: DeWitt Ross & Stevens, S.C.
                                                                                                          ó
                                                                                                          Gaps
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                                                                                                          ;
0
                                                                   Length 37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 38;
                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATCHTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                              0.5%; Score 26.8; DB 1;
illarity 93.3%; Pred. No. 9.1e+02;
Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.5%; Score 26.8; DB 1;
93.3%; Pred. No. 9.3e+02;
ative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: DeWitt Ross & Stevens, S.C.
8000 Excelsior Drive, Suite 401
                                                                                                                                              2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2310 CACACACAGACAGACACACACACACACA 2339
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APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTONEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 198, Application US/08222177A Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; MOLECULE TYPE: DNA (genomic); IMMEDIATE SOURCE; ; CLONE: mfd6rs
US-08-222-177A-68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Weber, James L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 38 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wisconsin
                                                          Query Match
Best Local Similarity
Matches 28; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 8000 E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                       US-08-222-177A-140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-222-177A-198
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28;
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Sequence 137, Application US/08222177A

Patent No. 5582979

GENERAL INFORMATION:
APPLICANT: Weber. James L.
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460

CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                            ö
                                                                                                   Query Match 0.5%; Score 26.8; DB 1; Length 39; Best Local Similarity 93.3%; Pred. No. 9.4e+02; Matches 28; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.5%; Score 26.8; DB 1; Length 39; Best Local Similarity 93.3%; Pred. No. 9.4e+02; Matches 28; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ); Db _
9.4e+02;
2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
FILING DATE:
                                                                                                                                                                                     2310 CACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2310 CACACACACAGACAGACACACACACACACA 2339
                                                                                                                                                                                                             8 CACACACACACACATACACACACACACACACA 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,40000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 CACACACACACACACACACACACACACACA 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 43
US-08-222-177A-119
; Sequence 119; Application US/08222177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; IMMEDIATE SOURCE:
; CLONE: mfd29rs
US-08-222-177A-137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 39 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Madison
STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                   IMMEDIATE SOURCE:
                                           ; CLONE: mfd12rs
US-08-222-177A-86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                            US-08-222-177A-137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
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                                                                                                                                                                                                                                                                                        RESULT 42
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APPLICANT: Weber, James L.
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 26.8; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2310 CACACACACAGACAGACACACACACACA 2339
               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT: INFORMATION:
NAME: Sara, Charles S.
RGSISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 0807/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
RESISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDUUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-222-177A-86
; Sequence 86, Application US/08222177A
; Patent No. 5582979
                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
JS-08-222-177A-397
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Best Local Similarity 93.3%;
Matches 28; Conservative (
                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 38 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 39 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Madison
STATE: Wisconsin
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FILING DATE:
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Gaps

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APPLICANT: Weber, James L. TITLE OF INVENTION: LENGTH POLYMORPHISMS IN TITLE OF INVENTION: (dG-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460
; Patent No. 5582979
; GRMERAL INFORMATION:
APPLICANT: Weber, James L.
ITILE OF INVENTION: LENGTH POLYMORPHISMS IN
ITILE OF INVENTION: (dG-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
INUMBER OF SEQUENCES: 460
; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                              ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2310 CACACACACAGACAGACACACACACACACA 2339
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                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
PRIOR APPLICATION DATE:
APPLICATION NUMBER: US
FILING DATE: 21-APR-1989
ATTORNES/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09865.601
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                    PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                        E: Floppy disk
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA (genomic);
iMMEDIATE SOURCE:
CLONE: mfd23rs
US-08-222-177A-119
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(608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
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                                                                                                                                                                                                              Wisconsin
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Best Local Similarity
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                                                                                                                                                                                        Madison
                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                               MEDIUM TYPE:
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US-08-222-177A-400
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                                                                                                                                                                                                                                                                                                                     COMPUTER:
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GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dc-dx)n.(dc-dx)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CUCREMY APPLICATION DATA:
APPLICATION NUMBER: US/08/222,117A
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 CACACACACACACACACACACACACACA 31
                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09865.601
                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
FILING DATE:
                                                                                                                                                                                                                                                                 09865.601
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 098
TELECOMMUNICATION INFORMATION
TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 091
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                     400:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear

MOLECULE TYPE: DNA (genomic)
US-08-222-177A-400
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 40 base pairs
                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 40 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Madison
STATE: Wisconsin
COUNTRY: USA
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ZIP: 53717-1914
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US-08-222-177A-152
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TITLE OF INVENTION: LENGTH POLYMORPHISMS IN TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME UNUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 238, Application US/08222177A
Patent No. 5582979
GENERAL INFORMATION:
TITLE OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
NUMBER OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dewitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                             COUNTRY: USA

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARRE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
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93.1%; Pred. No. 1.4e+03;
tive 0; Mismatches 2;
                                                                                                    ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2311 ACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERNCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09865.601
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IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 29 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
                                                                                                                                                                    Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wisconsin: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sest Local Similarity
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ZIP: 53717-1914
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , CLONE: mfdl0rs
US-08-222-177A-80
                                                                                                                                             Madison
                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-222-177A-238
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APPLICANT: Sergei M. Gryaznov
TITLE OF INVENTION: Convergent Synthesis of Branched and Multiply
TITLE OF INVENTION: Connected Macromolecular Structures
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooley Godward LLP
STREET: Five Palo Alto Square, 3000 El Camino Real
CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                          ö
                                                                                                                                                                        Query Match

0.5%; Score 26.2; DB 1; Length 40;
Best Local Similarity 79.5%; Pred. No. 1.3e+03;
Matches 31; Conservative 0; Mismatches 8; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.5%; Score 25.8; DB 1; Length 29; 93.1%; Pred. No. 1.4e+03; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                          2311 ACACACACAGACACACACACACACAACAATTAACA 2349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/455,627 FILING DATE: 31-MAY-1995
                                                                                                                                                                                                                                                                                   1 ACACACACACACACACACACACACACACACACACACA 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: NAKABULA, Jackie N.
REGISTRATION NUMBER: 35,966
REFERENCE/DOCKET NUMBER: LYXX-003/01 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-843-5000
TELEFAX: 415-857-0663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2310 CACACACACAGACAGACACACACACAC 2338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 80, Application US/08222177A Patent No. 5582979 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25, Application US/08455627 Patent No. 5571677
                                         TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfd34rs
US-08-222-177A-152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Weber, James L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 93.18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 29 nucleotides
nucleic acid
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STRANDEDNESS: sing]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94306-2155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 3 CLASSIFICATION:
                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                   RESULT 46
US-08-455-627-25
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US-08-222-177A-373

Sequence 373, Application US/08222177A

Sequence 373, Application US/08222177A

Patent No. 5582979

GENERAL INFORMATION:
APPLICANT: Weber, James L.
TITLE OF INVENTION: (dc-da)n.(dg-dT)n SEQUENCES AND METHODS OF USING SAME NUMBER OF SEQUENCES: 460

CORRESPONDENCE ADDRESS:
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0.5%: Score 25.8; DB 1; Length 30;
Best Local Similarity 93.1%; Pred. No. 1.4e+03;
Matches 27; Conservative 0; Mismatches 2; Indels
                                                                              Length 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/222,177A
                                                                           Query Match
0.5%; Score 25.8; DB 2;
Best Local Similarity 93.1%; Pred. No. 1.4e+03;
Matches 27; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: DeWitt Ross & Stevens, S.C. STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2311 ACACACACAGACAGACACACACACACA 2339
                                                                                                                                                                   CLASSIFICATION: 435
CLASSIFICATION TATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY_AGENT INFORMATION:
NAME: Sara, Charles S.
RECISTRATION NUMBER: 30,492
RECISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELEPHONE: (608) 831-2100
TELEFAX: (608) 831-2106
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA (genomic); IMMEDIATE SOURCE:
CLONE: mfd118rs
US-08-222-177A-373
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nucleic acid
EDNESS: double
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wisconsin
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                       MOLECULE TYPE: DNA
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 53717-1914
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Patent No. 583068

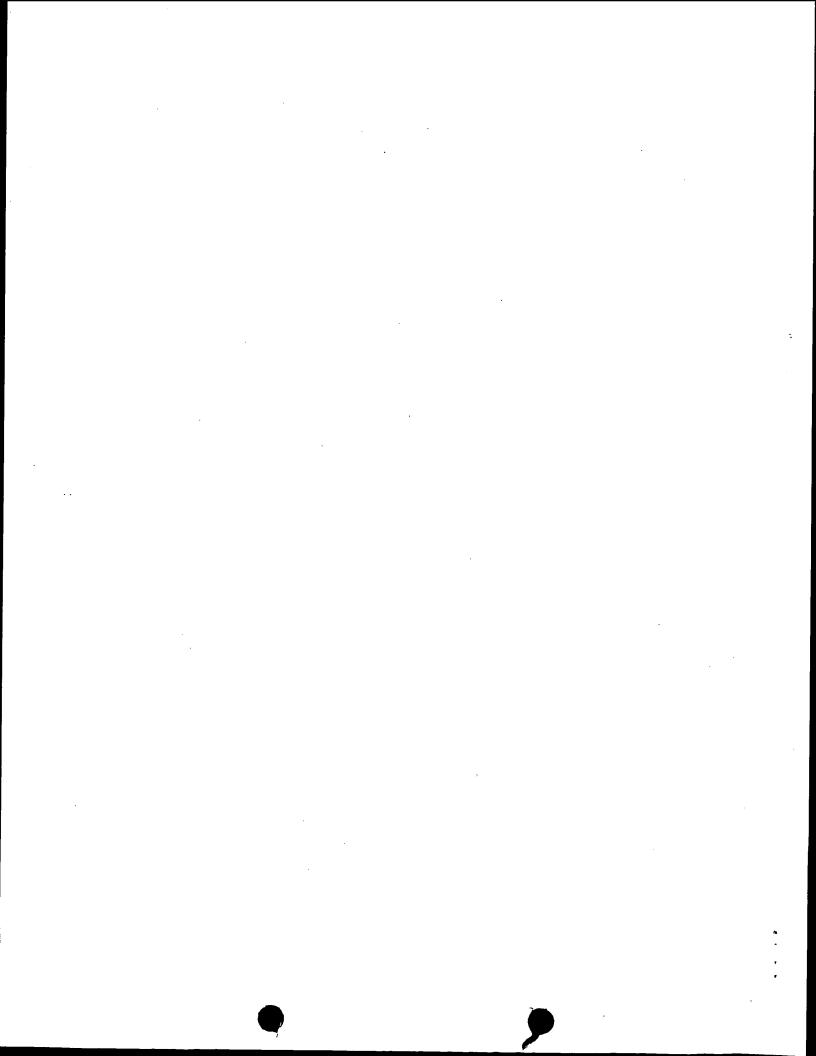
GENERAL INFORMATION:
TITLE OF INVENTION: Convergent Synthesis of Branched and Multiply TITLE OF INVENTION: Connected Macromolecular Structures NUMBER OF SEQUENCES: 26

CORRESPONDENCE ADDRESS:
ADDRESSE: COOLEY GOWARD LLP
STREET: Five Palo Alto Square, 3000 El Camino Real
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.5%; Score 25.8; DB 1; Length 29; 93.1%; Pred. No. 1.4e+03; tive 0; Mismatches 2; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/689,856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Nakamura, Jackie N.
REGISTRATION NUMBER: 35,966
REFERENCE/DOCKET NUMBER: LYNX-003/01 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-843-5000
TELEPHONE: 415-843-5000
INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2310 CACACACAGACAGACACACACACACACAC 2338
    FILING DATE:
CLASSIPICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: SATE, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 30,492
TELECOMUNICATION INFORMATION:
TELECHORE: (608) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09865.601
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APPLICATION NUMBER: US/08/455,627
FILING DATE: 31-MAY-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 238: SEQUENCE CHARACTERISTICS: LENGTH: 29 base pairs TYPE: nucleic acid STRANDEDNESS: double
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MOLECULE TYPE: DNA (genomic)
IMMEDIATE SOURCE:
CLONE: mfd65rs
US-08-222-177A-238
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TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity 93.1%
Matches 27; Conservative
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Gaps



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Copyright (c) 1993 - 2003 Compugen Ltd.
                                                            Total number of hits satisfying chosen parameters:
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               using sw model
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Maximum Match 100%
Listing first 500 &
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AZ827502 2M0103020 AZ827502 2M0103020 AZ316355 1M0034H10 AZ3164538 1M0034H10 AZ764538 1M056NN21 AUZ58912 AUZ69499 AZ404047 1M0112A10 AZ48464 1M0311H13 AZ620923 1M0458917 AZ793186 2M0468913 AZ48464 1M0311H13 AZ620923 1M0089K13 AZ753189 2M0008708 BZ894682 60143992 AZ456524 1M0259118 AZ456524 1M0259118 AZ456524 1M0259118 AZ677209 2M0003708 BZ894682 601439924 AZ351309 1M0696712 AUZ671278 AUZ71278 BZ774099 2M0003708 BZ894682 602294924 AZ35687 1M00565D2 AZ35687 1M00565D2 AZ35681 1M0152124 BZ356910 AUZ66910 AZ386218 1M0145M04 AZ66910 AUZ66910 AZ386218 1M0145M04 AZ609583 1M0560M05 AZ609583 1M0560M05 AZ784499 1M0560M22	AZ811930 ZM011021 AZ88198 M0164H14 AZ861607 ZM0164H14 AZ861607 ZM0164F11 AZ967442 ZM0238F20 AZ969917 ZM0242D20 AL752534 Arabidops B1078346 602874206 AZ970687 ZM024123 AA934710 0072a03.s AL587823 AL587823 AZ623276 1M0460M16
	0.4 35 17 A2831936 0.4 35 17 A2851899 0.4 35 17 A2951899 0.4 35 17 A296917 0.4 35 17 A296917 0.4 35 17 A2970697 0.4 36 17 A2970687 0.4 37 9 A39710 0.4 37 9 A39710 0.4 37 17 A2623276 0.4 37 17 A2623276 0.4 37 17 A2623276 0.4 38 10 AW3398 0.4 38 11 AL763877 13889 139 E ALIGNN AL
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AZ829195 ZMO106M12 AZ647335 IMO513415 AZ766498 IMO564E08 AZ861612 ZMO168P16 AL013688 HMO5 Sapi AU013658 AV0713658 AV674152 AV674152 AW334249 S3214 AGS AZ34541 IMO495F14 AZ6845779 ZMO145813 AL687908 tp89912.x AL68790 IMO47622 AZ774991 ZMO04722 AZ774991 ZMO04722 AZ627849 IMO74102 BF138797 GO2036229 AZ31115 IMO04102 BH000551 ZMO288F03 BH000551 ZMO288F03 BH76113 GO20495159 AZ44222 IMO236D07 BF525501 GO2069592 BF526501 GO20384505 AZ486773 1M0311B02 AZ486773 2M0010D15 AZ486773 2M0010D15 AZ486773 2M003731995 AZ789792 ZM001731	AZ626965 IN0467E15 AZ763999 IN0559208 AZ66737 IN0559208 AZ4825156 ZM0100N08 AZ418224 IN0186702 AZ486010 IN0313715 AL480085 T. brucel AI67781 wc80c06.x AZ486010 IN0313715 AZ486763 IN031815 AZ486763 IN031805 AZ51671 IN0250105 AZ58671 IN0250105 AZ58671 IN0250105 AZ661578 IN054028 AZ661578 IN054028 AZ661578 IN054028 AZ66157 W086617 AZ66157 W086617 AZ328537 IN0052823 AL484122 T. brucel AZ728538 IN0550317 AL744437 Danio rer AZ600409 IN0418407 AL03149 Homo sapi DZ5655 HUMGS04003 AL037510 HOMO sapi AZ683703 AL638703 AZ68209 IN0347E10 AZ586209 IN0347E10 AZ586209 IN0347E10
23 17 AZ829195 25 17 AZ766498 31 17 AZ766498 32 2 HSM00156 33 9 AU013658 34 10 AW674152 34 10 AW374249 34 10 AW374249 35 17 AZ686745 35 17 AZ686745 35 17 AZ686745 35 17 AZ686745 35 17 AZ686745 36 17 AZ774981 27 17 AZ774981 27 17 AZ774981 27 17 AZ774981 27 17 AZ774981 27 17 AZ774981 27 17 AZ774981 28 17 AZ774981 36 17 AZ774981 37 10 AW698832 35 17 AZ774981 36 17 AZ774981 37 17 AZ774981 38 12 AZ7818797 38 12 AZ7818793 39 17 AZ442523 38 17 AZ442523 39 17 AZ442523 39 17 AZ442613 39 17 AZ444253 39 17 AZ444253	11777117771777777777777777777777777777
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1 (bases 1 to 37)
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          REFERENCE
                                AUTHORS
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KEYWORDS
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                                                                                                                              TITLE
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                                                                                                                                                               Contact: Robert B. Weiss
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Wm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
84112, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (http://www.jdx.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polymerlectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil473114|gb|AR129072.1), a copy number inducible derivative of plasmid R1. The vector was ligated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XLI0-Gold (Stratagene) cells and selected for ampicillin resistance."
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1M0514J14F Mouse 10kb plasmid UUGCIM library Mus musculus genomic
                      Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus G57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                               Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 31, DB 17; Length 39;
Pred. No. 8.5e+04;
0; Mismatches 5; Indels
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                                                                                                                                                                                                                                                                                                                                                                                       Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                           Plate: 0081 row: B column: 23
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  High quality sequence stop: 39.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone="UUGC2M0081B23"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /strain="C57BL/6J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /sex="Male"
       (bases 1 to 39)
                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 87.2
Matches 34; Conservative
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AZ647581/c
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(http://www.jax.org/resources/documents/dhares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gild/32114 gp) half129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chamically-competent E. coli XI10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                               Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly iM., Rose, M., Rose, M., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/G7 (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                        Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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/clone="UUGCIMO514J14"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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Seq primer: CGTTGTAAAACGACGGCCAGT
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                                                                                                                                                                                                        Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Best Local Similarity 91.2%;
Matches 31; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
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//note="Vector: PWD42DV; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarcse gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114) [plant29072.1], a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the linsert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 36)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
JM., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
                                                                                                                                                                                                             Unpublished (2000)
Contact: Robert B. Weiss
Contract: Y of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /clone_lib="Mouse 10kb plasmid UUGClM library"
/sex="Male"
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                   Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0032 row: K column: 19
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 36.
Location/Qualifiers
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/clone="UUGC1M0032K19"
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Fax: 801 585 7177
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                                      AUTHORS
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AZ346800
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                                                                                                                                                                                                   Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Wan 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
84112, USA
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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1M0032K19R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0032K19 R, DNA sequence.
      1 (bases 1 to 35)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Dunn, D., Longarce, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
.M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
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/lab.host="E. Coli strain XLIO-Gold, Tl-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                                             Mouse whole genome scaffolding with paired end reads from 10kb
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/clone_lib="Mouse 10kb plasmid UUGClM library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Insert Length: 10000 Std Error: Plate: 0113 row: N column: 06 Seq primer: CGTTGTAAAACGACGCCAGT class: plasmid ends
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/db_xref="taxon:10090"
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                             plasmid inserts
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AZ315626/c
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                          Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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AZ304764.1 GI:10341104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Class: plasmid ends
            1 (bases 1 to 39)
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                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
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                                      AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMD42 (gil4732114) plant29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                            plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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1 (bases 1 to 37)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Slam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
JM., Rose,M., Rose,M., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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MO150D06F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0150D06 F, DNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                    Plate: 0082 row: 0 column: 09
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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/clone="UUGC1M0082009"
                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /sex="Male"
                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                          TITLE
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gal electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gil4732114[gb]AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                        Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /lab_host="E. Coli strain XLIO-Gold, T1-resistant, F-" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
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84.2%; Pred. No. 3e+05;
ative 0; Mismatches 6;
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                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0150 row: D column: 06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Mus musculus"
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/clone="UUGC1M0150D06"
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Location/Qualifiers
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REFERENCE AUTHORS

JOURNAL

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TITLE

FEATURES

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// AlaD. Most—E. Coll Strain Allo Colu, it resistant, it is musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with 74 DNA polymerase and 74
polymucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (qi|4732114|qbl|AR129072.1), a copy-number
inducible derivative of plasmid Rl. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XLIO-Gold (Stratagene) cells
and selected for ampicillin resistance."
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Mus musculus
Eukaryota; Hetazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Aammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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1M0498B20R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
                   Dunn, D., Acyagi A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                          Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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                                                                                                                                             Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /strain="C57BL/6J"
/db_xref="taxon:10090"
/db_xref="taxon:1000064L05"
/clone_lib="Mouse 10kb plasmid UUGClM library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0064 row: L column: 05
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Class: plasmid ends
High quality sequence stop: 38.
Location/Qualifiers
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1 (bases 1 to 38)
                                                                                                                                                                                                             Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                plasmid inserts
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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2M0064L05F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0064L05 F, DNA sequence.
   1 (bases 1 to 34)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Longarce, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
J.M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                             Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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/clone="UUGCIMO005A14"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
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0.6%; Score 28.2; DB 17; Length 34;
Best Local Similarity 90.9%; Pred. No. 3.4e+05;
Matches 30; Conservative 0; Mismatches 3; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0005 row: A column: 14
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Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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Fax: 801 585 7177
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                         Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Slam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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1M0112B09F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
                                                                                                                                                                                             Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                        Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /db_xref="taxon:10090"
/db_xref="taxon:10090"
/clone="UUGC1M0498B20"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.5%; Score 27.8; DB 17; Length 33; Best Local Similarity 93.5%; Pred. No. 4.1e+05;
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                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: C
Plate: 0498 row: B column: 20
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Location/Qualifiers
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                                                                                                                                                 plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
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                            AUTHORS
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114[gb]AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and
                                                                                                                                                                                                                                Unpublished (2000)
Contact: RObert B. Weiss
Contact: Robert B. Weiss
University of Utah
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 34)

Unun,D., Aoyagi,A., Barber,W., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Stokes,R., Tingey,A., von Niederhausern,A.,
and Wright,D.,Weiss,R.
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1M0439P13R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                                                        Mouse whole genome scaffolding with paired end reads from 10kb
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Pred. No. 4.1e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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Seq primer: CGTTGTAAAACGACGGCCAGT
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/db_xref="taxon:10090"
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AZ612777.1 GI:11734967
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us-09-375-514-19.rst

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/strain="C57BL/6J"
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Best Local Similarity 93.5%;
Matches 29; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /sex="Male"
1 (bases 1 to 35)
                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
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AZ819087/c
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                                                                                                                                            plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                       Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Slam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Brror: 0.00
Plate: 0439 row: P column: 13
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Mus musculus"
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/clone="UUGC1M0439P13"
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AZ626427.1 GI:11748617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex="Male"
  1 (bases 1 to 35)
                                                                                                                                                                                                                                                                                                                       Tel: 801 585 5606
Fax: 801 585 7177
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AZ626427
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VERSION
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarcse gel electrophoresis. Vector DNA was prepared from a derivative of pWM2 (gil4732114|gb|AF129072.1), a copy.number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli X110-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                    plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutherla; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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ZM0089G02R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGc2M0089G02 R, DNA sequence.
AZ819087
AZ819087.1 GI:12988995
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. Moneo, E., Welss, R.
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                                                                                                                                                   Mouse whole genome scaffolding with paired end reads from 10kb
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Pred. No. 4.1e+05;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0466 row: K column: 09
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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    .35
    /organism="Mus musculus"

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Location/Qualifiers
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/clone="UUGC1M0466K09"
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114) gblAR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the linsert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected for amplcillin resistance."
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 36) Dun, D., Acyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longarce, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="Vector: PWD42nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                      Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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                                                                                                                                    Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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                                                                                                                                                                                                Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Location/Qualifiers
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USA
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                                                                                                                                                                                                Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                           Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly M., Rose, M., Rose, M., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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1M0184D09F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
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/lab_host="E. Coli strain XLIO-Gold, T1-resistant, F-"
                                                                                                                                             Mouse whole genome scaffolding with paired end reads from 10kb
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Location/Qualifiers
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   (bases 1 to 35)
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AZ411170/c
LOCUS
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http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and 74 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwMo42 (gil4732114[gb]AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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1M0562C19R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0562C19 R, DNA sequence.
                        Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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                                                                                                                                            Mouse whole genome scaffolding with paired end reads from 10kb
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/strain="C5/BL/6J"
/db_xrref="taxon:10090"
/clone="UUGCLM0081M12"
/clone=lib="Mouse 10kb plasmid UUGClM library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Location/Qualifiers
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AZ765740.1 GI:12882075
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(bases 1 to 39)
                                                                                                                                                                                                                                                                                                                                                                                  rel: 801 585 5606
                                                                                                                                                                                                                                                                                                                                                                                                             Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                 Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT 84112, USA
121, USA
Tel: 801 585 506
Fax: 801 585 7177
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae, Mus.
                        Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Stan, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly iM., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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/note-"Vector: PMA2nv; Purified genomic DNA from M.
musculus C57BL/67 (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                         Mouse whole genome scaffolding with paired end reads from 10kb
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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Insert Length: 10000 Std Error: 0.00
Plate: 0101 row: F column: 02
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Class: plasmid ends
High quality sequence stop: 37.
Location/Qualifiers
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    /organism="Mus musculus"
    /strain="C57BL/6J"

                                                                                                                                                                                       Unpublished (2000)
Contact: Robert B. Weiss
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Rese
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   (bases 1 to 37)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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1 (bases 1 to 39)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Blan,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
.M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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0; Mismatches
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Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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                                                                                                                                                                                                               plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
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Best Local Similarity 93.55
Matches 29; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
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AUTHORS
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch oritice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMDA2 (gill4f92114fgplAR129072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
1 (bases 1 to 40)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacres,S., Mahmoud,M., Menen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R. Stokes,R., Tingey,A., von Niederhausern,A. Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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Seq primer: CACACAGGAAACAGCTATGACC
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/strain="C57BL/6J"
/db_xref="taxon:10090"
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AZ606329.1 GI:11728519
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Matches 29; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dhares/). The DNA was bydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114/1951AFL29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                      Unpublished (2000)
Contact: Robert B. Weiss
University of Utah
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
              1 (bases 1 to 33)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Lalam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
'M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
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                                                                                                                                                     Mouse whole genome scaffolding with paired end reads from 10kb
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
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                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0117 row: N column: 01
Seq primer: CGTTCTAAACGACGGCCAGT
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/strain="C57BL/6J"
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High quality sequence stop: 33.
Location/Qualifiers
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AZ373001.1 GI:10486701
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Fax: 801 585 7177
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SOURCE
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AZ373001/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/60 (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polymucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into
                                                                                                                                                                                                                                                                                                                    Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT 84112, USA
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Mus muscula
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 32)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
M., Ross,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Welss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
plasmid inserts.
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Pred. No. 5.5e+05;
0; Mismatches 3; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Plate: 0428 row: K column: 07
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculus"
                                                                                                                                                                                                      Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/clone="UUGC1M0428K07"
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Location/Qualifiers
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AZ834388.1 GI:13004296
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Best Local Similarity 90.69
                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                               Email:
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AZ834388/c
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Gaps

Std Error: 0.00

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Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
  1 (bases 1 to 36)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Slam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
'M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0517 row: E column: 02
Seg primer: CACACAGGAAACAGCTAYGACC
                                                                                                                                                                                                                                                                                                                       Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                USA
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AZ848143/c
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                                                                                                                                                                                          Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
84112, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10.5 Kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of PMD42 (gil4732114)gblAR129072.1), a copyrnumber inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                      Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly M., Rose, M., Rose, M., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note-"Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                       Mouse whole genome scaffolding with paired end reads from 10kb
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     High quality sequence stop: 36.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
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                                                                                                                                                                   Unpublished (2000)
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Matches 29; Conservative
                                                                                                                                                                                                                                                                                                               Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                              plasmid inserts
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (qi|q/32114|qb|aff22072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chantically-competent E. coli Xil0-Gold (Stratagene) cells and selected for ampicillin resistance."
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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2M0149A17F Mouse 10kb plasmid UGC1M library Mus musculus genomic
                                                                                                                                                                                                                                                                                                                                                                     /note="Vector: PWD42nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                     /lab_host="E. Coli strain XL10-Gold, Tl-resistant, F-"
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/clone_llb="Mouse 10kb plasmid UUGC1M library"
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90.6%; Pred. No. 5.5e+05;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    clone UUGC2M0149A17 F, DNA sequence.
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                                                                                                                               /organism="Mus musculus"
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                            High quality sequence stop: 36.
Location/Qualifiers
                                                                                                                                                                                               /db_xref="taxon:10090"
                                                                                                                                                               /strain="C57BL/6J
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                                                                                                                                                                                                                                                                                                         /sex="Male"
Class: plasmid ends
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Best Local Similarity
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1 (bases 1 to 39)

Junal, Adyad, Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly,
M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.

Mand Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
plasmid inserts
                                                                                                                                                                                                                                                       Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Matches 29; Conservative
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Fax: 801 585 7177
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AZ387867/c
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AUTHORS
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
   1 (bases 1 to 36)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
'M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
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                                                                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0149 row: A column: 17
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Seg primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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/db_xref="taxon:10090"
/clone="UUGC2M0149A17"
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and 74 polymerlectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (qilq1921HqiplaFA22072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into
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Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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1M0147N21R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0147N21 R, DNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /clone="UUGC2M0210G04"
/clone_lib="Mouse 10kb plasmid UUGC2M library"
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Pred. No. 5.4e+05;
0; Mismatches 3; Indels 0
                                                                                                                                                            Std Error: 0.00
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                                                                                                                                                            Insert Length; 10000 Std Error: (Plate: 0210 row: G column: 04 Seg primer: CACACAGAAACAGCTATGACC Class: plasmid ends

    .39
    /organism="Mus musculus"

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                                                                                                                                                                                                                                                                                                                                                                                                                                                        /db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                              High quality sequence stop: 39.
Location/Qualifiers
                                                                                                                                 Email: ddunn@genetics.utah.edu
                                                                                                                                                                                                                                                                                                                                                                                                                            /strain-"C57BL/6J
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /sex="Female"
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                           Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
                                                                                                                                                  plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
       1 (bases 1 to 36)
Dunn, D., Aoyagi, A.,
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                    Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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                                                                                                                                                                                          Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
84112, USA
Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                    Mouse whole genome scaffolding with paired end reads from 10kb
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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Pred. No. 5.4e+05;
0; Mismatches 8; Indels (
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                                                                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0147 row: N column: 21
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
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Location/Qualifiers
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                                                                                                                                                                    Unpublished (2000)
  1 (bases 1 to 40)
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                                                                                                                                               plasmid inserts
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Matches 32; Conserv
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                                                                                                                    TITLE
                                                                                                                                                                                          COMMENT
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (qi|4732114|qib|AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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1M0215L08R Mouse 10kb plasmid UUGCLM library Mus musculus genomic
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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/db_xref="taxon:10090"
/clone="UGGC1M0367L06"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                Std Error: 0.00
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                                                                                                                                       Plate: 0367 row: L column: 06
Seg primer: CGTTGTAAAACGACGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                 /organism="Mus musculus"
                                                                                                                                                                                                                                                        High quality sequence stop: 36.
                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
                                                                                                                                                                                                                                                                                              Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /sex="Male"
Tel: 801 585 5606
Fax: 801 585 7177
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//note="Vector: PWAZIVY; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was bydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gelectrophoresis. Vector DNA was prepared from a derivative of pWA2 (qil4732114 ipblARI29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for amplcillin resistance."
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Musinae; Mus.
1 (bases 1 to 40)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Slam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
i.M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                      308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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                                                                                                                                            Mouse whole genome scaffolding with paired end reads from 10kb
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/sex="Male"
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0005 row: B column: 05
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                                                                                                                                                                                                                              Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/clone="UUGC1M0005B05"
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                                                                                                                                                                                                      Unpublished (2000)
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Fax: 801 585 7177
                                                                                                                                                                          plasmid inserts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inco orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and 74 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gil473214[gb]AR129072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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1M0005B05R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0005B05 R, DNA sequence.
      1 (bases 1 to 37)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
AM., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /lab_host="E. Coli strain XLIO-Gold, Tl-resistant, F-"
/note="Vector: PWA21vr; Purified genomic DNA from M.
musculus G57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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0.5%; Score 27; DB 17; Length 37;
Best Local Similarity 85.7%; Pred. No. 6e+05;
Matches 30; Conservative 0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0215 row: L column: 08
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Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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/strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                                                                                                                                            Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
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Best Local Similarity 50.00
Matches 28; Conservative
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Fax: 801 585 7177
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                       Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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/note="Vector: PWP42nv; Purified genomic DNA from M.
musculus G57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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0; Mismatches 2.
                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0294 row: P column: 15
Seq primer: CGTGTAAAACGACGCCAGT
Class: plasmid ends
High quality sequence stop: 30.
Location/Qualifiers
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                                                                                                                                                                     Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/db_xref="taxon:10090"
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     (bases 1 to 30)
                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orlifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114) [pl]ARL29072.1], a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampleillin resistance."
                                                                                                                                                                                                                                        University of Utah

Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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2M0230G12R Mouse 10kb plasmid UUGC2M library Mus musculus genomic
1 (bases 1 to 30)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="E. Coli strain XLIO-Gold, T1-resistant, F-" /note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Std Error: 0.00
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Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: ddunn@genetics.utah.edu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AZ961995.1 GI:13833222
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1 (bases 1 to 31)
                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
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                   REFERENCE
                                       AUTHORS
                                                                                                                                                                                     JOURNAL
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SOURCE
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                                                                                                                                        TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /Jab_host="E" coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus G75BL/60 (female) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polymorlectide kinase. Adaptor oligonuclectides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (q14732114 lpb]AR129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                            Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT 84112, USA
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l (bases 1 to 30)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
'M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                            Mouse whole genome scaffolding with paired end reads from 10kb
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0230G12"
/clone_lib="mouse 10kb plasmid UUGC2M library"
/sex="Female"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ch 0.5%; Score 26.8; DB 17; Length 30; 1 Similarity 93.3%; Pred. No. 6.8e+05; 28; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                     Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0230 row: G column: 12
Seq primer: CACACAGGAACAGGTATGACC
Class: plasmid ends
High quality sequence stop: 30.
Location/Qualifiers
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                                                                                                                                                                           Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
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            REFERENCE
                                  AUTHORS
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pmAp2 (gql147221141gblAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XII0-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                          Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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1M0284N13R Mouse 10kb plasmid UGGCLM library Mus musculus genomic
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, M., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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/lab_host="E. Coli strain Xil0-Gold, Tl-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus G57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                                             Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0263 row: 0 column: 12
Seq primer: CACACAGGAAACAGCTATGACC
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    .31
    /organism="Mus musculus"

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Class: plasmid ends
High quality sequence stop: 31.
Location/Qualifiers
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/clone="UUGC1M0263012"
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AZ470636.1 GI:10628761
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1 (bases 1 to 31)
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  REFERENCE
                          AUTHORS
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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31 bp DNA linear GSS 04-OCT-200
1M0300N01R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0300N01 R, DNA sequence.
     1 (bases 1 to 31)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Longarce, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
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                                                                                                                         Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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Seq primer: CACACAGGAAACAGGTATGACC
Class: plasmid ends ends estop: 31.
High quality sequence stop: 31.
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/db_xref="taxon:10090"
/clone="UUGC1M0284N13"
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Fax: 801 585 7177
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AZ479719/C
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(http://www.jax.orgyresources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114) iplAFL29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for amplialin resistance."
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Dunn, D., Aoyaqi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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/clone_lib="Mouse 10kb plasmid UUGCIM library"
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Insert Length: 10000 Std Error: 0.00
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Seq primer: CACACAGGAAACAGCTATGACC
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                                                                                                                                                                                               Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Location/Qualifiers
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/db_xref="taxon:10090"
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Plate: 0300 row: N c
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                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
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1 (bases 1 to 31)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longarce, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
'M., Rose, M., Rose, R.,
and Wright, D., Weiss, R.,
Mouse whole genome scaffolding with paired end reads from 10kb
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Fax: 801 585 7177
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1 (bases 1 to 31)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
'M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                      Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0330 row: D column: 03
Seq primer: CACACAGGAAACAGCTATGACC
Class: plaamid ends
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Location/Qualifiers
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/clone="UUGC1M0330D03"
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AZ666438/C
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(http://www.jax.org/resources/documents/dhares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114)[pb]AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
plasmid inserts
Unpublished (2000)
Contact: Robert B. Welss
University of Utah
Gm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="E. Coli strain XLIO-Gold, Tl-resistant, F-"
/note="Weetcr: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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/clone_lib="Mouse_10kb plasmid_UUGCIM_library"
/sex="Male"
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                                                                                                                                                                                                                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0548 row: N column: 21
Seq primer: CGTTGTAAAACGACGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"/strain="C57BL/6J"
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Location/Qualifiers
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and 74 polymerase and 75 polymerase an
                                                                                                                                                                                                                                                                                                                                                                             Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
bunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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/db_xref="taxon:10090"
/clone="MGC2M018BC07"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0188 row: C column: 07
Seg primer: CGTTGTAAAACGACGCCAGT
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                                                                                                                                                                                                                                       Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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              REFERENCE
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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2M0188C07F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0188C07 F, DNA sequence.
1 (bases 1 to 31)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
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/db_xref="taxon:10090"
/clone="UGGIM0570M12"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0570 row: M column: 12
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Seq primer: CACACAGGAAACAGCTATGACC
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                                                                                                                                                                                                                                                     Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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High quality sequence stop: 31.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
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Matches 28; Conserv
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                     REFERENCE
                                                      AUTHORS
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                    2M0269D16R Mouse 10kb plasmid UUGC2M library Mus musculus genomic clone UUGC2M0269D16 R, DNA sequence.
                                                 Gaps
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0.5%; Score 26.8; DB 17; Length 31; 93.3%; Pred. No. 6.7e+05; tive 0; Mismatches 2; Indels (
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high .molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114) ppl.ARL29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for amplcillin resistance."
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Unpublished (2000)
Unpublished (2000)
University of Utah Genome Center
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            1 (bases 1 to 32)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Longarcer, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
i.M., Rose, M., Rose, R.,
and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /strain="C57BL/64"
/db_xref="taxon:10090"
/db_crone="UGC1M010A03"
/clone="UGC1M010A03"
/sex="Male"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0010 row: A column: 03
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Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Location/Qualifiers
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AZ311840.1 GI:10355201
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Matches 28; Conservative
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Fax: 801 585 7177
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                                                                                                                                                                                                                                                                                                                                                                                                                           Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT 121, USA 125 5606
Tel: 801 585 5606
Email: 601 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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1 (bases 1 to 31)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
J.M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Weiss, R.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                                                                                    Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/db_xref="taxon:10090"
/clone="UUGC2M0269DL6"
/clone="lib="Mouse 10kb plasmid UUGC2M library"
/sex="Female"
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Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Location/Qualifiers
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Matches 28; Conserv
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AUTHORS
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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  1 (bases 1 to 32)
Dun,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
.M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
                                                                                                                          Mouse whole genome scaffolding with paired end reads from 10kb
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/db_xref="taxon:10090"
/clone="MGCIMO27M1"
/clone="ID-"Mouse 10kb plasmid UUGCIM library"
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                                                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
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Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"
                                                                                                                                                       plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114 fighlAFL29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into
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Sukaryotas Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. Monse whole whele when and Witght, D., Weiss, R.
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                                                                                                                                                  Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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0.5%; Score 26.8; DB 17; Length 32;
Best Local Similarity 93.3%; Pred. No. 6.7e+05;
Matches 28; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.
Plate: 0132 row: F column: 15
Seq primer: CGTrGTAAAACGACGGCCAGT
Class: plasmid ends
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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                 Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rosse, M., Rosse, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Welss, R. Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts.
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32 bp DNA linear GSS 05-OCT-20C
1M0331E05F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0331E05 F, DNA sequence.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0246 row: M column: 11
Seq primer: CGTTGTAAAACGACGCCAGT
Class: plasmid ends
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                                                                                                                                                            Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/db_xref="taxon:10090"
/clone="UUGC1M0246M11"
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    (bases 1 to 32)
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                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA.polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114)fplARI29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Stokes, R., Tingey, A., von Niederhausern, A.
Mouse Wight, D., Weiss, R.
Mouse wight, D., Weiss, R.
Mouse wight and Wrible genome scaffolding with paired end reads from 10kb plasmid inserts
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                                                                                                                                                      Unpublished (2000)
Contact: Robert B. Weiss
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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1M0273P11R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0273P11 R, DNA sequence.
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/clone_lib="Mouse 10kb plasmid UUGCIM library"
/sex="Male"
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Best Local Similarity 93.3%; Pred. No. 6.7e
Matches 28; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0331 row: E column: 05
Seq primer: CGTTGTAAAACGACGGCCAGT
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/strain="C57BL/6J"
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Location/Qualifiers
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Mus musculus
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                                                                                                                                                                                Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
WR. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
84112, USA
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                       Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly M., Rose, M., Rose, M., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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1M0406C21R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0406C21 R, DNA sequence.
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                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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/clone="UUGCIM0273P11"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
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                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0273 row: P column: 11
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 33.
Location/Qualifiers
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                                                                                                                                                                   Unpublished (2000)
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Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
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// Arte="Nector: PWAZIV: Further Garonic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gil4732114/gb]AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
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1 (bases 1 to 33)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
, M., Rose, M., Stokes, R., Tingey, A., von Niederhausern, A.
and Witght, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
plasmid inserts
Unbullished (200)
Contact: Robert B. Weiss
University of Utah
Genome Center
University of Utah
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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MA0461G04R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0461G04 R, DNA sequence.
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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93.3%; Pred. No. 6.7e+05;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0406 row: C column: 21
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /strain="C57BL/6J"
/db_xref="taxon:10090"
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Fax: 801 585 7177
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// Alabinost Fig. Coll Strain All Colu, it resistant, it musculus C57BL/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (qi|4732114|qbb|AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
bunn,D., Aoyagi,A., Barber,W., Beacorn,T., Duval,B., Hamil,C.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.,
Mand Wright,D.,Weiss,R.
                                                                                                                                                                                                                                               Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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/sex="Male"
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
                                                                                                                                        Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0461 row: G column: 24
Seg primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
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/organism="Mus musculus"
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/clone="UUGC1M0461G24"
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Location/Qualifiers
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                                                                                                                                                                                                                   Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
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                                           AUTHORS
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Weiss B. Dinmedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
84112, USA
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
         1 (bases 1 to 33)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
ISlan, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly
IN. Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A.
and Wright, D., Welss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
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33 bp DNA linear GSS 13-DEC-200
1M0461G24R Mouse_10kb plasmid UGC1M library Mus musculus genomic
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/clone_lib-"Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0461 row: G column: 04
Seg primer: CACACAGGAAACAGCTATGACC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
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Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
          Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Stokes,R., Tingey,A., von Niederhausern,A., M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A., M., C., L., Welss,R.
                                                                                                                                         Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                            plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
        1 (bases 1 to 33)
Dunn, D., Aoyagi, A.,
                                                                                                                                                                                                                                                                                                                                                 Tel: 801 585 5606
Fax: 801 585 7177
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                                                                                                                                                                       L Unpublished (2000)
Contact: Robert B. Weiss
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah Genome Center
University of Utah
Rm. 308, Blomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT
84,112 USA
Fax: 801 585 5606
Fax: 801 585 7177
Email: doun@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0087 row: P column: 19
Seq primer: GGTTGTAAAACGACGCCAGT
Class: plasmid ends
High quality sequence stop: 33.
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          Loases 1 to 33)

Dunn,D., Aoyadi,J. Barber,M., Beacorn,T., Duval,B., Hamil,C.,
Islam,B., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
'M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
Mouse whole genome scaffolding with paired end reads from 10kb
plasmid inserts
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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2M0156015F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC2M0156015 F, DNA sequence.
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/clone="UUGC2M0087P19"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
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Pred. No. 6.7e+05;
0; Mismatches 2; Indels (
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Best Local Similarity
Matches 28; Conserv
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and 74 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gilg73211), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coll XL10-Gold (Stratagene) cells and selected for amplcillin resistance."
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Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0156 row: 0 column: 15
Seq primer: CGTTGTAAAACGACGGCCAGT
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/clone="UUGC2M0156015"
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Location/Qualifiers
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Matches 28; Conservative
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